OM nucleic - nucleic search, using sw model Run on:

November 19, 2002, 00:33:44; Search time 1026.35 Seconds (without alignments) 765.605 Million cell updates/sec

US-09-829-004A-9 27 1 aaacttcagtgtgtggacctccatgtt 27 Perfect score: Sequence:

IDENTITY_NUC Gapop 10.0 , Gapext 1.0 Scoring table:

2054640 segs, 14551402878 residues Searched:

4109280 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

gb_sts:*
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em_htg_other:* em_htg_hum:* em_htg_mus:* em_htg_pln:*

em_htg_mam:* em_htg_vrt:* em_htg_rod:*

em_htgo_mus:*
em_htgo_other:* em_htgo_hum:*

Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	AX281252 Sequence	3 Hum	342 Sequen	Metho		Ношо	luman p	edneuce	_	986 Sequenc	Sedu	M21895 Human prost	HOMO S	167864 Sequence 14	E32812 Primer DNA	X05332 Human mRNA	Homo	AJ310938 Homo sapi	M21896 Human prost	AR059540 Sequence	AR082934 Sequence	AX337560 Sequence	X07730 Human mRNA	AF335478 Homo sapi	AX106218 Sequence	AX033404 Sequence	AX033402 Sequence	AX033400 Sequence	Segu	Нишап	Human pro	യ	2	Human p	73 HOMO	502 Homo	13527 HOMO	0 M.mulatt	389 Canis	5 PSA-pro	3937 Homo sap	5710 Homo sap	11174 Homo sa	I19503 Sequence 1	351 Sequence	
CTTVUTTOC			HSPSA4	AR153342	BD007601	HSA459783	HSA459782 .	HSU17040	167863	AF335477	~	AX267642	•	HUMAPS	167864	E32812	HSPSAR	7	HSA310938	HUMPAB	AR059540 .	AR082934	AX337560	HSPSA	AF335478	AX106218	AX033404	AX033402	AX033400	AX033401	HSPSAG	HUMPSANTIG	A37262	AR167395	HUMPSAA	25	AC027602	2		AC090889	'n	601	AC016710	117	119503	167851	
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ALIGNMENTS

PAT 02-NOV-2001

linear

DNA

AX281252 27 bp Sequence 9 from Patent WO0176622. GI:16608508 synthetic construct. synthetic construct artificial sequences. AXŽ81252 AX281252.1 RESULT 1
AX281252
LOCUS
DEFINITION
ACCESSION
VERSION
KEYWORDS
SOURCE ORGANISM

Pedyczak,A., Chong,P. and Sia,C.D. Immunogenic peptides derived from prostate-specific antigen (psa) and uses thereof patent: WO 0176622-A 9 18-OCT-2001; REFERENCE AUTHORS TITLE JOURNAL

PAT 08-AUG-2001

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Gaps

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PAT 31-JAN-2002

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1. 711
/organism='Unidentified'
Location/Qualifiers
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Unclassified.
3 1 (bases 1 to 711)
5 Young/C.Y.F., Tindall, D.J. and Klee, G.G.
Nethod for detection of breast cancer
AL Patent: US 6235486-A 4 22-MAY-2001;
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Location/Qualifiers
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Submitted (09-JAN-1989) Digby M.R., The Howard Florey Institute,
Melbourne University, Parkville, Victoria 3052, Australia
2 (base 1 to 281)
2 (base 1 to 281)
2 (base 2 to 281)
2 (base 3 to 281)
2 (base 3 to 281)
3 (base 3 to 281)
4 theman prostate specific antigen (PSA) gene: structure and linkage Not the kallikrein-like gene, hGR-1
Nucleic Acids Res. 17 (5), 2137 (1989)
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Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammaliai; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Digby, M.R.
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Best Local Similarity 100.0%; Pred. No. 0.011;
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/note="prostate specific antigen"
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Aventis Pasteur Limited (CA)
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AJ459782.1 GI:20520640 "... alternative splicing; KLK3 gene; prostate specific antigen
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EPEEFLTPRKLQCVDLHYISNDVCAQVHPQKVTKFMLCAGRWTGGKSTCSGDSGGPLV
CNGVLQGITSWGSEPCALPERPSLYTKVVHYRKWIKDTIVANP"
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                                                                                                                                      HSA459783 870 bp mRNA linear PRI 09-MAY-2002
Homo sapiens mRNA for prostate specific antigen (KLK3 gene), splice
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                             Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire d'Enzymologie, Universite F. Rabelais, 2 bis byd Tonnelle, Tours,
                                                                                                                                                                                                                                                                                                     Heuze-Vourc'h,N. and Courty,Y. Complex alternative splicing of the hKLK3 gene coding for the tumour marker PSA (prostate-specific-antigen)
    Gaps
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Alternative splicing; KLK3 gene; prostate specific antigen.
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/translation-"MWVPVVPLTLSVTWIGAAPLILSRIVGGWECEKHSQPWQVLVAS
REARAVGGGVTRPPWPLTAAHGIRNESVILLGRISLEHPBPTOGQVFQVSHSFPPHLYD
MSLLKNRFLRPRDSSIEPBEFLTRKKLQVDLHVISNDVGAQVHPQKVTRENCAGR
WTGGKSKTGSOBGGPLVCNGVLQGITSWGSBPCALPBERPSLTTKVVHYRKWIKDTIVA
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A, complete cds.
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1 (bases 1 to 990)
Monne,M.M., Moreno,J.M., Mele,C.M., Mulholland,G.M. and
Gomella,L.G.
A Comparative Analysis of Prostate Specific Antigen Gene Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                          Euteleostomi;
                                                                                                                                                                                                                                                                                                         Tours,
                                                                                               Heuze-Vourc'h,N. and Courty,Y. Complex alternative splicing of the hKLK3 gene coding for the tumour marker PSA (prostate-specific-antigen)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                    Direct Submission
Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
d'Enzymologie, Universite F. Rabelais, 2 bis bvd Tonnelle,
37032 cedex, FRANCE
alternative splice variant sequences: AJ459783, AJ459784
                     Craniata; Vertebrata; Eutele
Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human prostate specific antigen precursor mRNA, U17040 U17040.1 GI:595945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /product="prostate specific antigen"
/protein_id="CAD30844.1"
/db_xref="G1:20520641"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /product="prostate specific antigen"
a 235 c 231 g 190 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note="alternative splice variant 1"
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100.0%; Score 27; DB 9;
Best Local Similarity 100.0%; Pred. No. 0.012;
Matches 27; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
1. 876
/organism="Homo sapiens".
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /evidence=experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9606"
/clone="YC171105-00"
Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /codon_start=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22. .72
/gene="KLK3"
94. .681
/gene="KLK3"
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22. .684
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                                                                                                                                                                                                    2 (bases 1 to 876)
Courty, Y.
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PRI 13-MAY-2002
A, complete
                                                                                                                                                                                                                                                    Homo sapiens.

Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota, Metazoa; Chordata; Catarrhini; Hominidae; Botteleostomi; Libasa; Louiso; L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product="prostate-specific antigen variant 1"
/protein_id="AAL/32041"
/db_xxef="G1:18478570"
/translation="MWVPVVFLTLSVTWIGERGHGWGDAGEGASPDCQAEALSPPTQH
PSPDRELGSFLSLPAPLQAHTPSPSILQQSSLPHQVPAPSHLPQNFLPIAQPAPCSQL
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David, A., Engel,S., Azar,I., Bernstein,J., Rotman,G., Savitsky,K.
and Mintz,L.
Direct Submission
Submitted (11-JAN-2001) Compugen Ltd., 72 Pinchas Rosen, Tel Aviv
69512, Israel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo saplens
Bukaryota's Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1350)
M. J., Dillon, D., C., Mitcham, J.L., Harlocker, S.L., Jiang, Y.,
Reed, S.G., Kalos, M.D., Fanger, G.R., Day, C.H., Retter, M.W.,
Compositions and methods for the therapy and diagnosis of prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                              AF335477

1130 bp mRNA linear PR1
HOMO sapiens prostate-specific antigen variant 1 mRNA,
AF335477
AF3354771 GI:18478569
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/codon_start=1
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1. 1130
Acronalism="Homo sapiens".
/db_xref="taxon:9606"
/chromosome="19"
/map="19913.3"
42. .356
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AX200986
Sequence 616 from Patent W00151633.
AX200986.1 GI:15390813
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Best Local Si
Matches 27;
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VERSION
KEYWORDS
SOURCE
ORGANISM
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DEFINITION
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AF335477
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AUTHORS
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AUTHORS
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ORGANISM
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AX200986
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                                                                                                                                                                                                                                        /organism="Homo sapiens"
/organism="Homo sapiens"
/organism="Igu"
/chzxef="taxon:9606"
/chromosome="19"
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/tissue_type="prostate specific antigen precursor"
/codon_start=1
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RGARACGGVLUPPOWYLTAAHCIRNSSVILLGRHSLEHPEDTGOVFOVSHSPHPLYD
MSLLKNRELRPCDDSSHDLMLILLSEBPRELDAWKWONDLPTORPHPLYD
EPERLIPRKLOGVDLHYISMUPCAQVHPOWYPKMLGAGRWGGSI
CNGVLGGITSWGGSDCGALPERPSLYTKVVHYRKWIKDILVANP"
in Benign and Malignant Prostate Tissue
1 (Dases 1 to 990)
Moreno, J. (Dases 1 to 990)
Moreno, J. (Dases 1 to 990)
Direct Submission
Submitted (08-NOV-1994) Jose M J G Moreno, Urology, Thomas
Jefferson University Hospital, 1025 Walnut Street, Philadelphia, PA
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Groce.C., Gomella,L., Mulholland,S.Grant., Moreno,J.G. and
Nucleic acid primers for detecting micrometastasis of prostate
cancer
Patent: US 5674682-A 13 07-0cr-1997;
Location/Qualifiers
1. 992
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Ouery Match

Best Local Similarity 100.0%; Score 27; DB 6; Length 992;

Matches 27; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                114. 824
/product="prostate specific antigen"
93. 113
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291 c 273 g 217 t
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167863.
167863.1 GI:2829985
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1. .990
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291 c 272 g
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Unknown.
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mat_peptide
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ORIGIN
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Matches
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167863
LOCUS
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AUTHORS
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/protein_id="AAA59995.1"
/brotein_id="AAA59995.1"
/db_xref="G1:189524"
/db_xref="G1:189524"
/db_xref="G00-119-695"
/db_xref="C00-119-695"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/note="AA 18. .24 propeptide; AA 69 glycosylated Asn
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Homo sapiens prostate-specific antigen mRNA, complete cds.
M26663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note="prostate specific antigen signal peptide'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 1415;
                                                                                                                                                                                                                                                            precursor"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /product="prostate specific antigen"
328 a 361 c 410 g 316 t
176 bp upstream of PvuII site.
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                                                                                                                                                                                                                                                            /note="prostate specific antigen
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Homo sapiens prostate gland cDNA to mRNA.
Homo sapiens
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Pred. No. 0.012;
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/organism="Homo sapiens"
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/map="19q13.3"
/clone="PSA-20"
                                                        1. .1415
/organism="Homo sapiens"
/db_xref="taxon:9606"
/map="19q13.3"
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/gene="APS"
/note="G00-119-695"
25. .810
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Kalos,M.D., Fanger,G.R., Retter,M.W., Stolk,J.A., Day,C.H.,
Vedvick,T.S., Carter,D., Li,S.X., Wang,A., Skeiky,Y.A., Hepler,W.T.
and Henderson,R.A.
Compositions and methods for the therapy and diagnosis of prostate
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end, clone PA 75.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Riegman, P.H., Klaassen, P., van der Korput, J.A., Romijn, J.C.
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Biochem. Biophys. Res. Commun. 155 (1), 181-188 (1988)
88326297
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I Similarity 100.0%; Pred. No. 0.012;
27; Conservative 0; Mismatches 0; Indels
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Human prostate specific antigen (PA) gene,
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CORIXA CORPORATION (US)
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Patent: WO 0151633-A 616 19-JUL-2001;
CORIXA CORPORATION (US)
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AX267642
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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        /codon_start=1
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/db_xref="GI=18864"
/db_xref="GI=18864"
/db_xref="GI=18969"
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RGRAVGGVLVHPQWVLTAAHCIRNKSVILLGRHSLFHPEDTGQVFQVSHSFPHPLYD
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MSELKRRKLQCVDLHYISNDVCAQVHPQKVTWFMLCARRGTGGKSTCSGDSGGFLV
CNGVLQGITSWGSEPCALPERPSLYTKVVHYRKMIKDTIVANP"
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Croce, C., Gomella, L., Mulholland, S.Grant., Moreno, J.G. and
Fischer, R.
Nucleic acid primers for detecting micrometastasis of prostate
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Best Local Similarity 100.0%; Pred. No. 0.012;
Matches 27; Conservative 0; Mismatches 0; Indels
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/note="G00-119-695"
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /organism="unknown"
374 c 422 g
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372 c 419 g
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/gene="APS"
/note="G00-119-695"
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/gene="APS"
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VERSION E32812.1 G118623942

KEWRORDS

Unidentified

ORRANISM unidentified

ORRANISM unidentified

ORRANISM unidentified

OCRANISM unidentified

Nakagawara, H.

TITLE

Primer DAS and method for detecting mRNA encoding prostate

JOURNAL

PATHORIS CHARGE OF 1200006996-A 5 07-MAR-2000:

COMMENT

OS Unidentified

ON-MAR-2000

PR 28-AGG-1998 JP 1998243419

PR 14 NROKAZU NNKAGAMARA

PR 28-AGG-1998 JP 1998243419

OC Strandedness: Single;

CC Topology: Linear;

CC Topology: Linear;

CC Topology: Linear;

CC Topology: Linear;

CG Topol
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BE838775 RC6-FT019 BE838775 RC6-FN011 BE771925 CM3-FT009

AA654296 nt83c06.s

ESI: *

Database

DB sed DB sed

Minimum I Maximum I

Perfect score:

Run on:

Sequence:

Scoring table:

Searched:

AA594946 no40c01.s BF679591 602154033 BF678378 602085809 AI732097 nc21g11.x BF964653 602267731 AI547309 PN001_AH_ BF679168 602153741

ww-bio.llni.gov/bbrp/image/image.html

T29518 EST82635 Hu BF588848 7112c03.x

AA657663 BF858362 BF858359 BE769440 T29518 BF588848

17777

100.0 100.0 100.0 100.0

27 27 27 27 27

Score

Result . Q

AA654548 nt59d11.s AA579039 nf34g07.s

AI926979 wo68d11.x AA654527 nt59a12.s

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27; Conservative
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BF858359/c
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ORIGIN
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AUTHORS
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KEYWORDS
SOURCE
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                                                                                                                                                                                                                                                                                                                                    /ask_marge="45 years old"
//ab_host="04108"
//ab
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 196)
Dias Neto. E., Garcia Crrea.R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Coldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
Simpson,A.J.
Shotgun sequencing of the human transcriptome with ORF expressed
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Fax: +55-11-270401
Fax: +55-11-270001
Fax: +55-11-270001
This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gephtml2.pl?tl=RC5&t2=RC5-FT0193-1100-012-G09_1ts1=2000-11-21&t4=1)
Seq primer: puo 18 forward
High quality sequence stop: 190.
Location/Qualifiers
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
2020268
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Trace considered overall poor quality
Seq primer: -folm15 fwd. ET from Amersham
High quality sequence stop: 1.
Location/Qualifiers
                                                                                                                        // 1. 189
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/clone="laAGE:1205002"
/clone=lib="NCI_CGAP_PF3"
/sex="Male"
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BF858362
BF858362.1 GI:12246106
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AUTHORS
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BF858362
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KEYWORDS
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1. .196
/organism="Homo sapiens"
/db_xrof="taxon.9600"
/clone_lib="FT0193"
/dev_stage="Adult"
/note="Crgan: prostate_tumor; Vector: pucl8; Site_1: SmaI;
Site_2: SmaI; A mini-library was:made by cloning products
derived from ORESTES PCR (U.S. Letters Patent application
No. 196.716 - Ludwig Institute for Cancer Research)
profiles into the pUC 18 vector. Reverse transcription of
tissue mRNA and cDNA amplification were performed under
low stringency conditions.
41 a 58 c 55 g 42 t
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

(bases 1 to 239)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,

Madl,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,

Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,

Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Simpson, A.J.
Shotgun sequencing of the human transcriptome with ORF expressed sequence tags
Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
20202663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                            Gaps
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0
                                                                                                                                                                                                                                                                                                                             100.0%; Score 27; DB 12; Length 196; 100.0%; Pred. No. 0.16;
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                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                            146 AAACTTCAGTGTGTGACCTCCATGTT 172
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Query Match Best Local

Matches

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BASE COUNT

VERSION KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS

DEFINITION

RESULT 4 BE769440

ACCESSION

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Humania; Eutheraca; Chordata; Craniata; Vertebrata; Euteleostomi; Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarhini; Hominidae; Homo.

1 (Dases 1 to 272)

2 Adams, M.D., Kerlavage, A.R., Fleischmann, R.D., Fuldner, R.A., Bult C.J., Lee, N., Kirkness, E.F., Weinstock, K.G., Gocayne, J.D., White, O., Sutton, G., Blake, J.A., Eardeon, R.C., Chiu, M.W., Clayton, R.A., Cline, R.T., Cotton, M.D., Earle-Hughes, J. Fitzchwan, S.M., Glodek, A., Gnehm, C.L., Hanna, M.C., Hedblom, E., Hinkled, P.S., Kelley, J.M., Moreno-Palanques, R.F., McDonald, L.A., Marmaros, S.M., Merrick, J.M., Moreno-Palanques, R.F., McDonald, L.A., Nguyen, D.T., Pellegrino, S.M., Phillips, C.A., Ryder, S.E., Scott, J.L., Saudek, D.M., Shirley, R., Bednarik, D.P., Ceo, L., Cepeda, M.A., Coleman, T.A., Clilins, E.-J., Dimke, D., Fenzie, A., Fischer, C., Hastings, G.A., He, W.-W., Kusch, C., Ji, H., Li, H., Meissner, P.S., Olsen, H., Raymond, L., Wei, Y.-F., Wing, J., Xu, C., Yu, G.-Li, Ruben, S.M., Dillon, P. J., Fannon, M. R., Rosen, C.A., Haseltine, W.A., Fields, C., Fraser, C.M. and
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                                                                                                       T29518 272 bp mRNA linear EST 06-SEP-1995 EST828353 Human Prostate gland Homo sapiens CDNA 5' end similar to antigen, prostate specific (HT:2350), mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Venter, J.C.
Initial Assessment of Human Gene Diversity and Expression Patterns
Based Upon 83 Million Basepairs of cDNA Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: tdbinfo@tdb.tigr.org
. Cor clone avallability, additional sequence and expression information related to this EST, please contact the TIGR Database (tdbinfo@tdb.tigr.org)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 272;
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/note="Organ: prostate gland"
81 c 71 g 56 t
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/organism="Homo sapiens"
/db_xref="ATCC (inhost):106527"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The Institute for Genomic Research
932 Clopper Rd, Gaithersburg, MD 20878
Tel: 3018699056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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Location/Qualifiers
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BF588848
BF588848.1 GI:11681172
                                                                                                                                                                                                                                       T29518.1 GI:611616
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Venter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fax: 3018699423
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Best Local Simi
Matches 27;
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BF588848/c
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DEFINITION
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This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/carcipts/gethtml2.pl7tl=&t2=PM1-FT0028-030
700-001-c02_l&t3=2000-07-03&t4=1)
Seq primer: puc 18 forward
High quality sequence stop: 241.

Location/Qualifiers

1. .241

Location/Qualifiers

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1 (bases 1 to 241)

Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R., Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F., Goldman, G.H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H., Brunstein, A., deoliveira, P.S., Bucher, P., Jongeneel, C.V., O'Hare, M.J., Soares, F., Brentani, R.R., Reis, L.F., de Souza, S.J. and
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research:
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Shotgun sequencing of the human transcriptome with ORF expressed
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                                                                                                       Length 239;
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                                                                                                   ch 100.0%; Score 27; DB 12; Similarity 100.0%; Pred. No. 0.17; 27; Conservative 0; Mismatches 0;
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+55-11-2707001
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Gaps

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BP856613

INM MR1-F70200-311000-001-b11 F70200 Homo sapiens CDNA, mRNA sequence.

BP856613.1 GI:12244370

BP856613.1 GI:12244370

BP856613.1 GI:12244370

BP856613.1 GI:12244370

BP856613.1 GI:12244370

BP856613.1 GI:12244370

INM man alian suplems

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; CD seas 1 to Aw W Jr. (Arrivallo, Arrivallo, Arr
                                                                                                                                                                                                                                                       /organism="Homo sapiens"
/db_xref="taxon.9606"
/db_xref="taxon.9606"
/clone=lib="MAGE:982382"
/clone_lib="MoI_GGAP_Prl2"
/sex="male"
/lab_bost="DH10B"
/note="Vector: pAMP10; mRNA made from metastatic prostate lession of the bone, CDNA made by oligo-dT priming.
Non-directionally cloned. Size-selected on agarose gel, average insert size 600 bp. Library made by D. Krizman, NIH."
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Fax: +55-11-2707001
Email: sainpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL
311000-001-blist=2000.10-318t4=1)
Seq primer: puc 18 forward
High quality sequence stop: 269.
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//db_xref="taxon:9606"
//db_xref="taxon:9606"
//dclone_lib="FT0200"
//dev_stage="Adult"
//note="Organ: prostate_tumor; Vector: pucl8; Site_1: Smal;
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Best Local Similarity 100.0%; Score 27; DB 9; Length 310;
Matches 27; Conservative 0; Mismatches 0; Indels 0; Gaps
                                                                 found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/lmage/lmage.html
losert Length: 426 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham.
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Ra Nc1-CcAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),

Tumor Gene Indea (1997)

Contact: Robert Strausberg, Ph.D.

Emall: Gapbbs: **Temath.nih.gov

Tissue Procurement: W. Douglas Figg, Ph.D., Paul H. Duray, M.D.,

CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Preparation: David B. Krizman, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
                                                         Burkaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 288)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute; Cancer Genome Anatomy Project (CGAP),
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MENA linear EST 05-AUG-1997

Similar to gb:W21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);

AA522842.1 GI:2263554
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1 Similarity 100.0%; Score 27; DB 12; Length 288;
27; Conservative 0; Mismatches 0; Indels (
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/note="Organ: prostate_tumor; Vector: puc18; Site_1: Smal; Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196, 716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: +55-11-2704922
Fax: +55-11-2707001
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=RC56t2=RC5-FT0193-
201100-012-H066t3=2000-11-208t4=1)
Seq primer: puc lB forward
High quality sequence strar: 48
High quality sequence stop: 329.
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mRNA sequence.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 329)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shotgun sequencing of the human transcriptome with ORF expressed
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100.0%; Pred. No. 0.19;
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     Mismatches
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dias Neto, E., Garcia Correa, R.,
                                                                                               AAACTICAGIGIGIGGACCICCAIGIT 176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.

Michael Emment: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.

Michael Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Conne distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Seq primer: -40ml3 fwd. ET from Amersham

High quality sequence stop: 298.

I. 328 Qualifiers

I. coation/Qualifiers
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Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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/db_xref="taxon:9606"
/clone="IMAGE:1205098"
/clone_llb="NCI_CGAP_Pr3"
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/lab_host="DH10B"
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Unpublished (1997)
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Prazz.1 = 7704922
Fax: +55-11-2707001
Email: asimpsoneludwig.org.br
Email: asimpsoneludwig.org.br
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=&t2=CM3-FT0097-300
60-225-Cll&t3=2000.06-30&t4-1)
Seq Primer: puo 18 forward
High quality sequence start: 25
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Locata.
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ISM Homo sapiens

Nor-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

Tumor Gene Index

Norbert Strausberg, Ph.D.

Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.

Contact: Robert Strausberg, Ph.D.

Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.

CONTA Library Preparation: David B. Krizman, Ph.D.

CONA Library Preparation: David B. Krizman, Ph.D.

CONA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
                                                                'M.J', Soares,F., brentanl,K.K., Rels,L.F., Ge SOGGE,S.C. w.S.
Simpson,A.J.
Shotgun sequencing of the human transcriptome with ORF expressed
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shailar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
AA228953.1 GI:1851772
                                           stein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                           Contact: Simpson A.J.G.
Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                / Match 100.0%; Score 27;· DB 12; Length 354; Local Similarity 100.0%; Pred. No. 0.19; Or Indels (Mismatches 0; Indels (
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Frail +55-11-2704922
Fax: +55-11-2707021
Fax: +55-11-2707001
Fax: +55-11-2707001
Frail: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL 800-011-F04613-2000-008-098414=1)
Seq primer: puc 18 forward
High quality sequence start: 8
High quality sequence start: 8
High quality sequence stop: 278.
Location/Qualifiers
1. 351
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Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Dias Neto, E., Garcia Correa R., Verjovski-Almeida,S., Briones,M.R., Godfman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.W.Y., Soares,F.F., Matsukuma,A., Baia,G.S., Simpson,D.H., M.J., Soares,F.F., Brunstein,A., Geoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Simpson,A.J., Soares,F.F., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                              Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                             Shorgun sequencing of the human transcriptome with ORF expressed sequence tags
Proc. Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
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Homo sapiens cDNA, mRNA sequence.
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Butaryote;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
Dias Netco, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R.,
Rogai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F.,
Goldman, G. H., Carvalho, A.F., Matsukuma, A., Baia, G.S., Simpson, D.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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27; Conservative 0; Mismatches 0; Indels (
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EC771925.1 GI:10225583
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Email: cgapbs-remail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html
Insert Length: Siz Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
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/lab_host="DillOB"
/lab_host="DillOB"
/lab_host="DillOB"
/note="Vector: pAMPIO; Site_1: Not1; Site_2: EcoRI; 1st
strand cDNA was primed with oligo(dT)17 on 50 ng of
DNAse-treated, total cellular RNA obtained from 5,000-10
,000 microdissected, histologically normal prostate
epithelial cells. Double-stranded cDNA was ligated to
EcoRI adaptors, 5 cycles of PCR applied to the cDNA with
an adaptor-specific primer, and the resulting PCR product
subcloned into pAMPIO by the UDG-cloning method (Life
Technologies). Average insert size is 600 bp. NOTE: Not
directionally cloned. This library was constructed by
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strand cDNA was primed with oligo(dT)17 on 50 ng of
DNAse-treated, total cellular RNA obtained from 5,000-10
/OO microdissected cells histologically determined to be
fully malignant prostate cancer cells. Double-stranded
cDNA was ligated to EcoRI adaptors, 5 cycles of PCR
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at 1007.s1 NCI_CGAP_Pr3 Homo sapiens cDNA clone IMAGE:1203925
similar to gb:M21895 PROSTATE SPECIFIC/ANTIGEN PRECURSOR (HUMAN);
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Unpublished (1997)
On Nov 12, 1997 this sequence version replaced gi:2559129.
Contact: Robert Strausberg, Ph.D.
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/organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                         directionally cloned.
David Krizman."
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AA635287
AA635287.1 GI:2619092
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.(
Matches 27; Conservative
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AA635287
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TITLE
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KEYWORDS
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Tumor Gene Index
Dupublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nin.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llni.gov/Dbrp/Ainage/Ainage.html
Insert Length: 648 Std Error: 0.00
Seq primer: -41ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                      /dev_stage="45 years old"
/dev_stage="45 years old"
/lab_host="PHIOB"
/note="vector: pAMPIO; Site_1: Not1; Site_2: EcoRI; Ist
strand cDNA was primed with oligo(dT)17 on 50 ng of
strand cDNA was primed with oligo(dT)17 on 50 ng of
bNAse-treated, total cellular RNA obtained from 5,000-10
,000 microfaissected, histologically normal prostate
epithelial cells. Double-stranded cDNA was ligated to
EcoRI adaptors, 5 cycles of PCR applied to the cDNA with
an adaptor-specific primer, and the resulting PCR product
subcloned into pAMPIO by the UDG-cloning method (Life
Technologies). Average insert size is 600 bp. NOTE: Not
directionally cloned. This library was constructed by
David Krizman."
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ANTIGEN PRECURSOR (HUMAN);,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 388) NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap. NCI-GGAP http://www.ncbi.nlm.nih.gov/ncicgap. National Cancer Institute, Cancer Genome Anatomy Project (GGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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the I.M.A.G.E. Consortium/LLNL at:
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                      www-bio.llnl.gov/bbrp/image/image.html
Seg primer: -28mija revi Er from Amersham
High quality sequence stop: 342.
Location/Qualifiers
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                                                                                                                                                  /organism="Homo sapiens"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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Location/Qualifiers
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found through
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AA228822/c
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                                                                                                          FEATURES
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applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
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88 a BASE COUNT ORIGIN

0; Gaps Query Match 100.0%; Score 27; DB 9; Length 398; Best Local Similarity 100.0%; Pred. No. 0.2; Matches 27; Conservative 0; Mismatches 0; Indels

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Search completed: November 19, 2002, 05:17:33 Job time : 1545.46 secs

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Sequence 616, App
Sequence 616, App
Sequence 598, App
Sequence 977, App
Sequence 5, Appli
Sequence 3, Appli
Sequence 1, Appli
Sequence 2, Appli
Sequence 280, App
Sequence 280, App
Sequence 34, Appli
Sequence 34, Appli
Sequence 603, Appli
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Sequence 153, App
Sequence 334, App
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111.898 Million cell updates/sec
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                                                                                                         November 19, 2002, 04:04:51; Search time 91.3846 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cgn2_6/prodata/1/pubpna/US07_PUBCOMB.seq:*
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GenCore version 5.1.3
Copyright (c) 1993 - 2002 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-759-143-616
US-09-759-143-616
US-09-822-827-616
US-09-822-827-977
US-09-755-100-3
US-09-755-100-3
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US-09-755-100-1
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US-09-759-143-334
                                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                        1 aaacttcagtgtgtggacctccatgtt 27
                                                                                                                                                                                                                                                                                                                 335578 seqs, 189365133 residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Published_Applications_NA:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                        - nucleic search, using sw model
                                                                                                                                                                                                                                                             IDENTITY_NUC Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                            Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                             US-09-829-004A-9
27
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Match Length DB
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Gaps

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Length 27; Indels

Query Match
Best Local Similarity 100.0%; Pred. No. 0.00059;
Matches 27; Conservative 0; Mismatches 0;

US-09-829-004A-9

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Sequence 616, Application US/09759143 Patent No. US20020022248A1 GENERAL INFORMATION:

US-09-759-143-616

Xu, Jiangchun Dillon, Davin C. Mitcham, Jennifer

APPLICANT: APPLICANT: APPLICANT:

| Patent No. US20020132976A1 | Patent No. US20020132976A1 |
| Patent No. US20020132976A1 |
| Patent No. US20020132976A1 |
| CAPERAL INFORMATION: Pele |
| APPLICANT: Chong, Pele |
| APPLICANT: Sia, Charles Dwo Yuan |
| TITLE OF INVENTION: Immunogenic Peptides Derived from Prostate-Specific Antigen (P TITLE OF INVENTION: Uses Thereof |
| FILE REPERENCE: 11014-24 |
| CURRENT APPLICATION NUMBER: US/09/829,004A |
| CURRENT FILING DATE: 2001-04-16 |
| PRIOR FILING DATE: 2000-04-10 |
| NUMBER OF SEQ ID NOS: 12 |
| SEQ ID NOS: 12 |
| SEQ ID NO |
| SE Sequence 334, App Sequence 698, App Sequence 698, App Sequence 698, App Sequence 378, App Sequence 1379, App Sequence 1379, App Sequence 13191, A Sequence 1321, App Sequence 1268, App Sequence 101, App Sequence 2307, App Sequence 11, App Sequence 1, App Sequence 1, App Sequence 1, App Sequence 2307, App Sequence 1, App Sequence 2307, App Sequence 1, App Sequence 2307, App Sequence 2012, App Sequence 2013, App Sequence 2013, App Sequence 2013, App Sequence 20142, App Seq 5, Appli 1, Appli 3, Appli 1, Appli 938, Ap 142, Ap 083, Ap 0 US-09-780-669-334 0 US-09-822-827-334 0 US-09-759-143-698 0 US-09-780-669-698 0 US-09-914-511-985 0 US-09-954-511-985 US-09-954-511-1379 US-09-954-511-1379 US-09-961-511-1379 US-09-961-511-1379 US-09-961-511-1379 US-09-961-511-1379 US-09-961-511-1379 US-09-961-511-1379 US-09-961-511-1379 US-09-961-1375-68 US-09-961-1375-6 US-09-961-1395-4 US-10-041-395-4 ALIGNMENTS TYPE: DNA ORGANISM: Artificial Sequence 100 100 100 100 100 100 100 100 OTHER INFORMATION: CLP316 365 407 16552 16552 305 453 3360 3360 2652 2997 3531 3878 10378 16. 16. FEATURE: LENGTH:

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US-09-969-708-598
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                                                                                                                                                     APPLICANT: mcneil, Patricia D.
APPLICANT: mcneil, Patricia D.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND FILE REPERBACE: 210121.427C24
CURRENT APPLICATION NUMBER: US/09/780,669
CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 616
                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match
Best Local Similarity 100.0%; Score 27; DB 10;
Best Local Similarity 100.0%; Pred. No. 0.0013;
Matches 27; Conservative 0; Mismatches; 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 616, Application US/09780669; Patent No. US20020051977A1; GENERAL INFORMATION:
Harlocker, Susan L.
Jiang, Yuquiert A.
Henderson, Robert A.
Kalos, Michael D.
Fanger, Gary R.
Stetter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
Li, Samuel
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Dillon, Davin C.
Mitcham, Jennifer L.
Harlocker, Susan L.
Jiang, Yuqui
Henderson, Robert A.
Kalos, Michael D.
Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
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Vedvick, Thomas S.
Carter, Darrick
Li, Samuel
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Hepler, William
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                                                                                                                                                                                                                                                                                                                                                        ; TYPE: DNA
; ORGANISM: Homo sapien
US-09-759-143-616
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; ORGANISM: Homo sapien
US-09-780-669-616
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APPLICANT:
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Sequence 598, Application US/09969708

Patent No. US30020102532A1

GENERAL INFORMATION:

APPLICANT: AUGUSTUS.

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign

TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign

TITLE OF INVENTION: USA09/969,708

FILE REPERENCE: 689290-70

CURRENT FILING DATE: 2000-10-03

PRIOR APPLICATION NUMBER: US/60/237,608

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

NUMBER OF SEQ ID NOS: 658

SEQ ID NO 598

LENGTH: 1729
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                                                                                                                                                                                                                                           US-09-822-827-616
US-09-822-827-616
Sequence 616, Application US/09822827
Patent No. US2020020081860A1
GENERAL INFORMATION: All OF USE OF UNCOUNTIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210.11.5.41
CURRENT FILING DATE: 2001-03-28
UNMARE OF SEQ ID NOS: 982
SEQ ID NO 616
SEQ ID NO 616
LENGTH.1350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .;
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100.0%; Score 27; DB 10;
Best Local Similarity 100.0%; Pred. No., 0.0013;
Matches 27; Conservative 0; Mismatches 0;
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100.0%; Score 27; DB 10;
Best Local Similarity 100.0%; Pred. No. 0.0014;
Matches 27; Conservative 0; Mismatches 0;
Query Match
100.0%; Score 27; DB 10;
Best Local Similarity 100.0%; Pred. No. 0.0013;
Matches 27; Conservative 0; Mismatches 0;
                                                                                             Oy 1 AAACTTCAGTGTGTGGACCTCCATGTT 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
CORGANISM: Homo sapien
US-09-822-827-616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: DNA
; ORGANISM: Homosapiens
US-09-969-708-598
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Gaps

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GENERAL INFORMATIONS:
GENERAL INFORMATIONS:
GENERAL INFORMATIONS:
TITLE OF INFORMATION NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES;
FILE REFERENCE: 2786-0156.
GURRENT FILING DATE: 2001-008
CURRENT FILING DATE: 1099-02-18
PRIOR PELLING DATE: 1999-02-18
PRIOR PELLING DATE: 1999-04-14
PRIOR PELLING DATE: 1999-04-14
PRIOR PELLING DATE: 1999-04-14
PRIOR PELLING DATE: 1999-04-14
SPRIOR PELLING DATE: 1999-08-11
NUMBER OF SEC ID NOS: 12
SOFTWARE: PATENTIN VEY: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence I. Application US/09755100
Patent No. US2002009189A1
GENERAL INFORMATION:
TUTLE OF INVENTIOR: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 1999-02-18
PRIOR PLING DATE: 1999-02-18
PRIOR PLING DATE: 1999-04-14
PRIOR PLING DATE: 1999-04-14
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-08-11
                                                                                                                                        Query Match
Best Local Similarity 100.0%; Pred. No.:0.0016;
Matches 27; Conservative 0; Mismatches 0; Indels 0
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SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
LENGTH: 4661
                                                TYPE: DNA
CORGANISM: Homo sapiens
US-09-755-100-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Homo sapiens
US-09-755-100-3
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                                                                                                                                                                                                                                                                                                                                                                       RESULT 9
US-09-755-100-3
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US-09-755-100-1
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     SEQ ID NO 5
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; Patent No. US20020081680A1
; GENERAL INFORMATION:
    APPLICANT: Xu, Jiangchun
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
    TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
; FILE REFERENCE: 210121.534C1
; CURRENT APPLICATION NUMBER: US/09/822,827
; CURRENT FILING DATE: 2001-03-28
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 980
; LENCTH: 2406
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Patent No. US20020099189A1
GENERAL INFORMATION:
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION NUMBER: US/09/755,100
PRIOR APPLICATION NUMBER: US/09/755,100
PRIOR PELLING DATE: 1999-02-18
PRIOR PELLING DATE: 1999-02-18
PRIOR PELLING DATE: 1999-04-14
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-08-11
            Sequence 977, Application US/09822827
Patent No. US20020081680A1
AERREAL INFORMATION:
APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
FITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.53421
CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NOS: 992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Db 1537 AAACTTCAGTGTGTGACCTCCATGTT 1563
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SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                           ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-822-827-977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
CORGANISM: Homo sapiens
US-09-822-827-980
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US-09-822-827-977
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LOCATION: 89, 92, 122, 123, 295, 320, 322, 352, 355, 361, 369, 375,
LOCATION: 89, 92, 122, 123, 295, 320, 322, 352, 355, 361, 369, 375,
LOCATION: 882, 884, 386, 389, 409, 411, 413, 414, 441, 442,
LOCATION: 448, 453, 461, 560, 510, 528, 532, 537, 546, 547, 563, 564,
LOCATION: 569, 577, 583, 587, 598, 599, 609, 613, 616, 621
OTHER INFORMATION: n = A,T,C or G
LOCATION: 622, 627, 648, 681
COTHER INFORMATION: n = A,T,C or G
US-09-919-580-34
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                                                                               Sequence 72, Application US/09919580

Patent No. US20020110832a1

GENERAL INFORMATION:
APPLICANT: With
APPLICANT: With
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: ADD DIAGNOSIS OF COLÓN CANCER
TITLE OF INVENTION: AND DIAGNOSIS OF COLÓN CANCER
TITLE OF INVENTION: AND DIAGNOSIS OF COLÓN CANCER
FILE REFERENCE: 2010.12.55

CURRENT APPLICANTON NUMBER: US/09/919,580

CURRENT FILING DATE: 2001-07-30

NUMBER OF SEQ ID NOS 934

SEQ ID NO 72

SEQ ID NO 72

LENGTH: 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 34, Application US/09919580
Sequence 34, Application US/09919580
Sequence 34, Application US/09919580
Sequence 34, Application
Sequence 34, Application:
Sequence 34, Application:
Sequence 34, Application:
APPLICANT: Pyle, Ruth
APPLICANT: Xu, Jiangchun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 21011.552
CURRENT APPLICATION NUMBER: US/09/919,580:
CURRENT FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 934
SOFTWARE: FastEEQ for Windows Version 4.0
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-919-580-72
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US-09-919-580-34/c
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US-09-969-708-280/c
US-09-969-708-280/c
Sequence 280, Application US/09969708
Fatent No. US20020102532a1
GENERAL INFORMATION:
APPLICANT: Augustus, Meena
TITLE OF INVENTION: Sets
FILE OF INVENTION: Sets
FILE REFERENCE: 689200-70
CURRENT FILING DATE: 2000-10-03
FRICH REPLICATION NUMBER: US/60/237,606
FRICH REPLICATION NUMBER: US/60/237,608
FRICH APPLICATION NUMBER: US/60/237,608
FRICH REPLICATION NUMBER: US/60/237,608
FRICH REPLICATION NUMBER: US/60/237,425
FRICH REPLICATION NUMBER: US/60/237,425
FRICH REPLICATION NUMBER: US/60/237,425
FRICH FILING DATE: 2000-10-03
NUMBER OF SEQ ID NOS: 658
SOFTWARE: PatentIn Version 3.0
SEQ ID NO 280
SED ID NO 280
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APPLICANT: SATIZKY, Kinneret et al.

TITLE OF INVENTUREN: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REPERENCE: 2786-0156p
CURRENT FLING DATE: 1090-02-18

PRIOR APPLICATION NUMBER: UL 128587

PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14

PRIOR FILING DATE: 1999-04-14

PRIOR FILING DATE: 1999-04-14

PRIOR FILING DATE: 1999-06-11

NUMBER OF SEQ ID NOS: 12

SEQ ID NO 2

LENGTH: 4661

TYPE: DAA

VORGANISM: HOMO SapienS

US-09-755-100-2
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Pred. No. 0.0017; Mismatches 0;
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                                                                                                                                                                                                              RESULT 11
US-09-755-100-2
; Sequence 2, Application US/09755100
; Patent No. US20020099189A1
; GENERAL INFORMATION:
   Best Local Similarity 100.0%; Promatches 27; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CSGANISM: Homo sapiens US-09-969-708-280
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72.6%; Score 19.6; DB 10; Length 1168;
Best Local Similarity 84.6%; Pred. No. 4;
Matches 22; Conservative 0; Mismatches 4; Indels 0;
US-09-925-300-603

Sequence 603, Application US/09925300

Patent No. US20020151681A1

GENERAL INFORMATION:

APPLICANT: Craig Rosen,

APPLICANT: Steve Ruben

TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies

FILE REFERENCE: PAIO.

CURRENT APPLICATION NUMBER: US/09/925,300

CURRENT FILING DATE: 2001-08-10

PRIOR PRILING DATE: 2000-03-08

PRIOR PELING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 1890

SOFTWARE: PATENTION NUMBER: 60/124,270

PRIOR FILING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 1890

SOFTWARE: PATENTION NUMBER: 60/124,270

PRIOR FILING DATE: 1999-03-12

NUMBER OF SEQ ID NOS: 1890

SEQ ID NO 603

LENGTH: 1168

LCATION: (1121)

OTHER INFORMATION: n equals a,t,g, or c

LCATION: (1122)

OTHER INFORMATION: n equals a,t,g, or c

LCATION: (1133)

COTHER INFORMATION: n equals a,t,g, or c

LCATION: (1133)

COTHER INFORMATION: n equals a,t,g, or c

LCATION: (1133)

COTHER INFORMATION: n equals a,t,g, or c

LCATION: (1153)

COTHER INFORMATION: n equals a,t,g, or c

LCATION: (1153)

COTHER INFORMATION: n equals a,t,g, or c
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Run on:	November 19,	2002,	November 19, 2002, 00:33:44; Search time 912.308 Seconds (without alignments) 765.605 Million cell updates/sec
Title: US-09-829-004A-7 Perfect Score: 24	US-09-829-004A-7	4A-7	

24	٠.	residues
ggttgtcttcctc	Sapext 1.0	14551402878
24 1 atgtgggtcccggttgtcttcctc 24	IDENTITY_NUC Gapext 1.0	2054640 seqs, 14551402878 residues
Periect score: 24 Sequence: 1	Scoring table: IDENTITY_NUC Gapop 10.0,	Searched:

4109280	
g chosen parameters:	
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Listing first 45 summaries	GenEmbl:*	gb_ba:*	qb htg:*	qb_in:*	*: mo qb	*: vo_dp	gb_pat:*	gp_ph:*	db_pl:*	gp_pr:*	gb_ro:*	gb_sts:*	dp-sy:*	*:un_qb	gb_v1:*	em ba:	ent_tun: "	em in:*	em mu:*	em_om: *	em_or:*	em_ov:*	em_pat:*	em_ph:*	em_p1:*	em_ro:*	em_sts:*	em_un:*		htg_	htg.	em_ntg_other:*	em_htg_mus:*	em_htg_pln:*	em_htg_rod:*	htg	htg.	3 y :	em_htgo_hum:*	em_htgo_mus:*	em_htgo_other:*
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Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description,	AX281250 Sequence	XI3940 Human PSA g	S75755 PSA=prostat	AJ459783 HOMO Sapi	AJ4597	167863	٠, ٠	M26663	.*		AX033403 Sequence	AX106218 Sequence			M24543 Human prost	A37262 Sequence 2 AR167395 Sequence	M27274 Human prost	ACULISZS HOMO SAPI ACU27602 Homo Sapi	AF243527 Homo sapt	AX200977 Sequence	AX267633 Sequence	AX434375 Sequence	AC016126 Homo sapi 282215 Human DNA s	AC026616 Homo sapi	AC115837 Homo sapi AC009629 Homo sapi	3086 Homo	1479 Emer	S		DNA linear PAT 02-NOV-2001), prostate-specific antigen (psa)	11.
SUMMARIES	AX281250		875755	HSA3109 HSA4597	HSA459782	167863		HUMAPS	I67864 E32812			6 AX106218				6 A37262 6 AR167395			AF24352	AX20097		AX43437	AC01612			9 AK093086 8 AF459639		ALIGNMENT	٠	24 bp from Patent WO0176622.			ces.	g, P. and Sia, C.D. des derived from	2-A 7 18-OCT-2001
y h Length	100.0 24		_								 _		 							- m	m 16		~ ~		~ ~	76.7 1939	v			AX281250 Sequence 7 from Parcon Parco	81250.1 GI:1660850	synthetic construct		Pedyczak, A., Chong, P. Immunogenic peptides	and uses thereot Patent: WO 017662
Result No. Score															-				32	07	35	13	38 18	87	41 18	C 43 18.4	18		T 1 250	LOCUS AX2 DEFINITION Seq		KEYWORDS . SOURCE SYD		AUTHORS Ped	and JOURNAL Pat
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L (bases 1 to 70)
Everhart, D.S., Kaylor, R.M. and McGrath, K.
Patterned binding of functionalized microspheres for optical diffraction-based biosensors
Patent: US 6221579-A 2 24-ARR-2001;
Location/Qualifiers
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1899 bp DNA linear Numan PSA gene for prostate specific antigen exon 1. x13940.

181340.1 GI:35722

XABILIkreln; prostate specific antigen; PSA gene. Homo sapiens.

Homo sapiens.
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Location/Qualifiers
                                                   1. 24
/organism="synthetic construct"
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1 a 7 c 9 t
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Sequence 2 from patent US 6221579.
AR147236
Aventis Pasteur Limited (CA)
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S/2/55 See 15-JUN-1995 PSA-prostate-specific antigen [human, breast cancer specimen, mRNA partial, 569 nt]
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FQVSHSFPHPLXDMSLLKNRFLRPGDDSSHDLMLLRLSEPAELTDAVKVWDLPTQEPA
IGTTCYASGWGSIBPEBELIPFKKLQCVDL"
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Monne, M., Croce, C.M., Yu, H. and Diamandis, E.P.
Molecular characterization of prostate-specific antigen messenger RNA expressed in breast tumors.
Cancer Res. 54 (24), 6344-6347 (1994)
9507906
GenBank staff at the National Library of Medicine created this
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Best Local Similarity 100.0%; Pred. No. 0.67;
Matches 24; Conservative 0; Mismatches 0; Indels
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273. .>318
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hotte="prostate-specific antigen"
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Eur. J. Blochem. 268 (16), 4408-4413 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
Submitted (11-APR-2001) Courty Y., Faculte de Medecine, EMI-U 0010,
laboratoire d'Enzymologie, 2 bis bd Tonnelle, 37032 Tours cedex,
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MSLLKNRFLRPGDDSSHDLMLLRLSEPAELTDAVKVMDLPTQEPALGTTCYASGWGSI
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens mRNA for putative preproPSA-RP2 (KLK3 gene),
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                                            9; Length 569;
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/note="alternative splicing, transcript 1"
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                                          Score 24; DB 9;
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1. 708
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             related accession number AJ310938.
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22...807
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EPEFELTPRKLQCVDLHVISNDVCAQVHPQKVTKFMLCAGRWTGGKSTCSGDSGGPLV
CNGVLQGITSWGSEPCALPERRSLYTKVVVYRKWIKDTIVANP"
                                                                                                                                                                                                                                                                                                        HSA459783 870 bp mRNA linear PRI 09-MAY-2002
Homo sapiens mRNA for prostate specific antigen (KLK3 gene), splice
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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alternative splicing; KLK3 gene; "prostate specific antigen.
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ch 100.0%; Score 24; DB 9; Length 708; 1 Similarity 100.0%; Pred. No. 0.67; 24; Conservative 0; Mismatches 0; Indels
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257 c 242 g 186 t
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                                                                                                             1 ATGIGGGTCCCGGTTGTCTTCCTC 24
                                                                                                                                                38 ATGTGGGTCCCGGTTGTCTTCCTC 61
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22. .807
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94. .804
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Courty, Y.
Direct Submission
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/ 2. 356
//note="PSA; KLK3; alternatively spliced"
//note="PSA; KLK3; alternatively spliced"
//note="PSA; KLK3; alternatively spliced"
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//product="AaL73204.1"
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//protein_id="AaL73204.1"
//product="AaL73204.1"
//pr
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Homo sapiens prostate-specific antigen mRNA, complete cds.
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prostate-specific antigen.
Homo sapiens prostate gland cDNA to mRNA.
Homo sapiens
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Submitted (11-JAN-2001) Compugen Ltd., 72 Pinchas Rosen, Tel Aviv
69512, Israel
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Henttu.P. and Vihko.P.
CDNA coding for the entire human prostate specific antigen shows coding for the entire human tissue kallikrein genes high homologies to the human tissue kallikrein genes Biochem. Biophys. Res. Commun. 160 (2), 903-910 (1989) 28246551 2470373
On Jan 5, 1995 this sequence version replaced gi:341511.
Ref [1] reports bases 1. 135 only.
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and Mintz,L.
Unusual alternative splicing within the human kallikrein and KLR3 gives rise to novel prostate-specific proteins J. Biol. Chem. 277 (20), 18084-18090 (2002)
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100.0%; Score 24; DB 9; Length 1130;
Best Local Similarity 100.0%; Pred. No. 0.66;
Matches 24; Conservative 0; Mismatches 0; Indels 0
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David,A., Engel,S., Azar,I., Bernstein,J.
and Mintz,L.
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1. .1446
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1.1130
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25. .810
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                                                                                                                                          HSA459784 1095 bp mRNA linear PRI 09-MAY-2002
Home sapiens mRNA for prostate specific antigen (KLK3 gene), splice
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Direct Submission

Submitted (03-MAY-2002) Courty Y., EMI-U-0010, Laboratoire

G'Enzymologie, Universite F. Rabelais, 2 bis bvd Tonnelle, Tours,

37032 cedex, FRANCE
alternative splice variant sequences: Aj459782, AJ459783.

Loation/Qualifiers

1. 01095
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Complex alternative splicing of the hKLK3 gene coding for the
tumour marker PSA (prostate-specific-antigen)
Unpublished
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AJ459784.1 GI:20520644
alternative splicing; KLK3 gene; prostate specific antigen.
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/db_xref="taxon:9606"
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AF335477.1 GI:18478569
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38. .352
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Courty, Y.
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HUMBAN MRNA for prostate specific autigen; signal

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Matches 24; Conservative 0; Mismatches 0; Indels 0; C
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AUTHORS
TITLE
JOURNAL
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COMMENT
FEATURES
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                                             /Godon_start=1
/Fordon_start=1
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RSGRAVGGGVLVBPOWTIFAHGTENSSYLLIGHSPHEDTGOVFOWSHSPPHPLYD
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RGGRATGGGGFLV
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167864 I GI:2829986
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Unclassified.

Unclassified.

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I (bases 1 to 1462)

RS Croce, C., Gomella, L., Mulholland, S. Grant., Moreno, J. G. and Fischer, R.

Nucleic acid primers for detecting micrometastasis of prostate cancer

NAL Patent: US 567482-A 14 07-0CT-1997;

Location/Qualifiers

Location/Qualifiers

Organism="unknown" 343 a 374 c 422 g 323 t
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residue"
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Best Local Similarity 100.0%; Score 24; DB 9; Length 1446;
Matches 24; Conservative 0; Mismatches 0; Indels (
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Best Local Similarity 100.0%; Score 24; DB 6; Length 1462;
Matches 24; Conservative 0; Mismatches 0; Indels (
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a 372 c 419 g 322 t
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Codon_grart=1

// Codon_grart=1
// Protein_id="CAA20947.1"
// Ab_xref="G1:35741"
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MSLLKNRFLRFCRODSSHDMLAREPERPLICSRIVGGWECEKHSOPWQVLVAS
RGRAVCGGTTSRCACPACHTON CAQUET SPEETING SEPETING SEP
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Search completed: November 19, 2002, 04:02:53 Job time: 920.641 secs

(OT920) NNAJ8 3DA9 2111

GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compus

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Result
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length: 2000000000
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1: /SIDSZ/gcgdata/gc2: /SIDSZ/gcgdata/gc3: /SIDSZ/gcgdata/gc4: /SIDSZ/gcgdata/gc5: /SIDSZ/gcgdata/gc6: /SIDSZ/gcgdata/gc6: /SIDSZ/gcgdata/gc6: /SIDSZ/gcgdata/gc6: /SIDSZ/gcgdata/gc7: /SIDSZ/gcgdata/gc
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1 atgtgggtcccggttgtcttcctc
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Maximum Match 100%
Listing first 45 summaries
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        AAD21361
AAT61707
AAT35293
ABN36013
AAA63193
AAA63192
ABL60747
AAT92974
AAT92974
ABV07911
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Prostate specific Prostatic specific Forward primer F1 Human spliced tran Nucleic acid used Prostate specific cDNA for prostate Forward primer F2 Human prostate exp
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	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11	10
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	22	22	22	22																	21	19	20	18	19	23	23	21	20	18	21	23	22	21	23	23
	AAK89250	AAL05893	AAL05891	AAL09557	AAT34621	AAC31348	ABK84795	ABK75499	ABL95315	AAH93851	AAS63944	ABV37834	AAT61701	AAA64168	AAA64167	AAA64169	AAA64171	AAF86958	AAT35868	AAA64170	AAA11711	AAV32496	AAX99847	AAT91055	AAV32497	ABV29201	ABV23473	AAC03740	AAX99846	AAT91054	AAZ47134	ABV25078	AAH41845	AAA61482	$\frac{\omega}{\omega}$	ABV07396
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ALIGNMENTS .

RESULT 1 AAD21361

AAD21361 standard; DNA;

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10-APR-2000; 2000US-195456P
                                                                                                                                                                                         Prostate-specific antigen; PSA; immunogénic peptide; immune response; therapy; cancer; prostate cancer; tumour metatasis; prophylaxis; ds.
        (AVET ) AVENTIS PASTEUR LTD
                                                                  18-OCT-2001
                                                                                                                                                                                                                           Prostate specific antigen (PSA) derived peptide, CLP313 encoding DNA.
                                                                                                                                                                                                                                                 28-JAN-2002 (first entry)
                                              10-APR-2001; 2001WO-CA00473
                                                                                      WO200176622-A2
                                                                                                                                    /*tag=
                                                                                                                 /product= "PSA derived peptide, CL
/note= "CDS does not include stop
                                                                                                                                                          Location/Qualifiers
                                                                                                        /partial
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RESULT 2
AAT61707
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The patent discloses immunogenic peptides of prostate-specific antigen (PSA) and nucleic acids encoding them. The peptides of the invention are used to prepare a medicament to elicit an immune response in an animal. They are used to treat cancer such as prostate cancer and tumour metastasis. They are also useful for prophylaxis, for preparing monoclonal or polyclonal antibodies, and in conventional techniques of immunology, molecular biology, cell biology and recombinant DNA technology. The present sequence is a DNA encoding PSA derived peptide, CLP313.
                   Claim 6; Page 38; 49pp; English.
                                       Treatment of benign prostatic hyperplasia or prostatic cancer using an anti-sense oligo:nucleotide targeted against prostate specific antigen or the probasin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT61707 standard; cDNA; 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 7; Page 28; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polypeptide, useful for treating comprises prostate-specific antigen
                                                                                                                Zamecnik PA;
                                                                                                                                                            20-SEP-1995;
                                                                                                                                                                                                           27-MAR-1997
                                                                                                                                                                                                                                  W09711172-AJ
                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                   Prostatic specific antigen antisense oligonucleotide.
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                                                                                                                                                                                   20-SEP-1996;
                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                 Human; prostatic specific antigen; probasin; rat; cancer;
hyperplasia; antisense oligonucleotide; ss.
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                                                                                         1997-202881/18
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DB; AAE13023.
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/note= "Phosphorothioated linkages"
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancers such as prostate cancer, derived peptide -
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novel method has been developed for treating a patient diagnosed

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC comprises administering to the patient a therapeutic amount of a CC comprises administering to the patient a therapeutic amount of a CC composition comprising an antisense oligonucleotide which selectively pridises to prostate specific antigen (PSA) gene or mRNA or to a CC probasin gene or mRNA sequence of the patient, where the antisense CC oligonucleotide inhibits expression of the sequence. The present conjugate of the patient, where the antisense CC sequence represents a specifically claimed antisense oligonucleotide against prostatic specific antigen. The antisense oligonucleotide can be used to effectively inhibit the growth of (and for the killing of) CC hyperplastic cells or cancerous cells or prostatic origin. Expression of the PSA and probasin genes is specific to prostatic cells so that the CC antisense oligonucleotide can be administered systemically, making it particularly useful in late stage prostatic cancer which has metasers and in which the cells have become resistant to oestrogen coranti-androgen therapy. The method can also be used in benign prostate hyperplasia, or early stage prostatic cancer, as an alternative for the correct radical procedures currently used, such as transurethral resection, coradical prostatectomy, or physical or chemical castration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
AAT35293-T35295 represent amplification primers for the prostate specific antigen (PSA). These primers were used to insert the CDNA for PSA into a BamHI-NotI-cleaved pVL1939 baculovirus transfer vector. The vector that was produced is ATCC VR-2496. The rPSA can produced by the vector can be used as an immunoassay calibrator. The immunoassays that use the rPSA are for the quantitative determination of PSA in a biological sample. In the assay, the test sample is combined with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 27 BP; 9 A; 7 C; 9 G; 2 T; 0 other;
                                                                                                                                                                                        Recombinant fragments -
                                                                                                                                                                                                                                                                                                                                                     28-APR-1995;
03-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT35293;
                                                                                                                                      Claim 3; Page 5; 19pp; English.
                                                                                                                                                                                                                                                WPI; 1996-356106/36.
                                                                                                                                                                                                                                                                                 Barnett TR,
                                                                                                                                                                                                                                                                                                                                                                                                          31-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                            07-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP725139-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Forward primer F1 for prostate specific antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT35293 standard; DNA;
                                                                                                                                                                          tests
                                                                                                                                                                                                                                                                                                                 (FARB ) BAYER CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   baculovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27
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24; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chain reaction; primer;
                                                                                                                                                                                        prostate-specific antigen (PSA) and immunological for use as an immunoassay calibrator in commercial PSA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transfer vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                               Ng PC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                     95US-0430498.
95US-0383111.
                                                                                                                                                                                                                                                                                                                                                                                                          96EP-0101304.
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                                                                                                                                                                                                                                                                               Yeung KK;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR; PSA; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           amplify; prostate specific
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cc (sub)transcriptome comprises messenger RNAs that populate a (sub)transcriptome, where the CC (sub)transcription units that populate a genome. The library comprises cCC several oligonucleotides, each capable of hybridising selectively to a Set of messenger RNAs transcribed from multiple CC the genome, which encodes one or more messenger RNA splice variants. The oligonucleotide libraries are useful for detecting mRNAs from a given transcription unit of CC quantitatively characterising the corresponding transcriptome, and in CC detecting RNA transcripts and splice variants of human or animal CC transcriptomes. The libraries may also be used as specialised mini CC libraries to detect transcripts of a sub-transcriptome under a CC particular biological or pathological state, and so allowing the CC condition; to detect developmental specific genes such as those genes conly expressed in specific tissue under a specific pathological CC transcripts and splice variants of a transcriptome under a specific so pathological state, and so allowing the CC condition; to detect developmental specific genes such as those genes conjugated and specific tissue under a specific pathological CC transcripts and splice variants of a transcriptome of a patient suffering CC transcripts and splice variants of a transcriptome of a patient suffering conjugated to detect developmental specific pathological conjugated to detect developmental specific pathological conjugated to detect developmental specific pathological conjugated to detect suffering conjugated to the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT ABN3601:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New oligonucleotide libraries comprising oligonucleotides which selectively hybridize to mRNAs transcribed from a transcription a genome, useful for detecting tissue-, pathology-, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shoshan A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JUL-2000; 2000US-221607P
02-MAY-2001; 2001US-287724P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; mouse; rat; splice transcript; detection; RNA transcriptione; oligonucleotide library;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABN36013 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                developmental-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-JUL-2001; 2001WO-IB01903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (COMP-) COMPUGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002-257383/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      spliced transcript detection oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID 8761; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Wasserman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 4 A; 15 C; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
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by performing the assay on one or more
an immunologically reactive fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВР
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Pred. No.
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0.16;
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Query Match

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Best Local
                                  analyte can be detected by analysing the diffiraction pattern of the transmitted light. The method of the present invention may used to detect and quantify a variety of analytes, e.g. midroparanisms, viruses, nucleic acids, hormones, steroids, antibodies, drugg, metals and antigens, either for clinical diagnosis of infections, cancer and allergies or for detecting microbial contamination of garments, The present sequence is a synthetic oligonucleotide, which was used, as an analyte to illustrate the
                                                                                                                             The present invention relates to a method for detecting and quantifying the presence or amount of an analyte within a medium. The method comprises micro-contact printing of analyte specific receptors onto polymer film. The target analyte binds to the polymer film, and results in light diffraction when light is transmitted through the film. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                \rm N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                  infections and cancer, comprises reacting analyte with diffraction-enhancing material carrying specific receptor
                                                                                                                                                                                                                                                           Diffraction-based analyte detection, useful for clinical diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Optical diffra contamination;
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                                                                                                                                                                                                           Example 2; Page 23; 38pp; English.
                                                                                                                                                                                                                                                                                                                                         (KIMB ) KIMBERLY-CLARK WORLDWIDE INC.
                                                                                                                                                                                                                                                                                                                                                                   11-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                              22-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                        15-JUN-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleic acid used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13 ATGTGGGTCCCGGTTGTCTTCCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ATGTGGGTCCCGGTTGTCTTCCTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                diffraction-based
nation; infection;
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                                                                                                                                                                                                                                                                                                                 Kaylor
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                                                                                                                                                                                                                                                                                                                                                                                              99WO-US27671.
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/mod_base= OTHER
/note= "Biotin-5'-"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Α,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       as an analyte in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        diffraction-based biosensor assay
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RESULT 6
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Matches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sig_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Prostate specific antigen; PSA; prostate; cancer; prostatic disease; gene; ds.
                    Sequence 90
                                           The invention provides a method of determining prostate specific antigen (pPSA) in a sample which involves determining the amount of complex formed on treating a sample with a pPSA specific antibody. The method is useful for detecting prostate cancer and distinguishing it from benign prostatic disease. The present sequence represents a DNA encoding a partial fragment of the PSA.
                                                                                                                                                   Determining prostate specific antibody (pPSA), comprises determining the amount of complex formed on treating a sample with a pPSA specific antibody, useful for detecting prostate cancer and distinguishing it from benign prostatic disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                      mat_peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prostate specific antigen (PSA) fragment encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL60747;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL60747 standard; DNA; 90
                                                                                                                             Disclosure; Fig 1; 30pp; English.
                                                                                                                                                                                                                                                  Mikolajczyk SD, Rittenhouse HG, Wang
                                                                                                                                                                                                                                                                                                                                   30-APR-1999;
                                                                                                                                                                                                                                                                                                                                                           23-FEB-2001; 2001US-0792534
                                                                                                                                                                                                                                                                                                                                                                                  18-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                          US2002045198-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-SEP-2002
                                                                                                                                                                                                                P-PSDB; ABB08071.
                                                                                                                                                                                                                                                                        (MIKO/) MIKOLAJCZYK S D.
(RITT/) RITTENHOUSE H G.
(WANG/) WANG T J.
(WOLF/) WOLFERT R L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002-425439/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 100.0%; 24; Conservative 0;
                     BP; 8 A; 26 C; 31 G; 25 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                   99US-0302965
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note=
52..72
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/note= "pa
1..72
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Propsa"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "preproPSA"
                                                                                                                                                                                                                                                                                                                                                                                                                                "partial mature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "partial fragment"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "PSA"
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; mismatches
                                                                                                                                                                                                                                                   ŢĮ,
                                                                                                                                                                                                                                                                                                                                                                                                                                 PSA"
                                                                                                                                                                                                                                                   Wolfert RL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA
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RESULT 8
AAT35294
ID AAT3
XX
AC AAT3
XX
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                                                                                                                δÃ
                                                                                             DЬ
                                                                                                                                          Query Match
Best Local S
Matches 24
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             AAT35294;
                                  AAT35294 standard; DNA; 93 BP
                                                                                            38 ATGTGGGTCCCGGTTGTCTTCCTC 61
                                                                                                                  1 ATGTGGGTCCCGGTTGTCTTCCTC 24
                                                                                                                                          l Similarity
24; Conserv
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Query Match

100.0%;

Score 24;

DB 24;

Length 90;

Conservative

0;

Mismatches

Indels

0;

Gaps

0

100.0%;

Score 24; Pred. No.

0.17;

DB 18; 0;

Length 92;

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cDNA sequences AAT92974-77 encode the prostate specific antigen (PSA). The detection of PSA mRNA in peripheral blood is associated with metastatic prostate cancer. Detection of PSA mRNA in the bloodstream indicates that prostate cells are circulating in the blood and confirms diagnosis of cancer. Circulating prostate cells also indicate a risk of a secondary tumour or bone metastatic anchoring. Detection of the PSA mRNA after prostatectomy or radiotherapy indicates a risk that the prostate cancer has spread and that the surgery and radiotherapy was not effective. PCR primers AAT92971-73 were used for the monitoring and diagnosis of patients with prostate cancer. The method comprises the detection of PSA mRNA by reverse transcriptase polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diagnosis; prostate cancer; prostate specific antigen; PSA; metastatic prostate cancer; prostate cell; secondary tumour; bone metastatic anchoring; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Monitoring or diagnosis of prostate cancer - by detecting prostate-specific antigen mRNA using specific primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA for prostate specific antigen (PSA).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-APR-1998 (first entry)
Sequence 92 BP; 14 A; 28 C; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 13; 25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Robbins DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT92974 standard; cDNA; 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SMIK ) SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9739139-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1997-526473/48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97WO-US06497
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   G; 22 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
      0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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RESULT 9
ABV07911
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                                                                                                                                                                                                                          Query Match
Best Local :
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                                                                                                                                                                                                                                                              AAT35293-T35295 represent amplification primers for the prostate specific antigen (PSA). These primers were used to insert the cDNA for PSA into a BamHI-NotI-cleaved pVLI393 baculovirus transfer vector. The vector that was produced is ATCC VR-2496. The rPSA can produced by the vector can be used as an immunoassay calibrator. The immunoassays that use the rPSA are for the quantitative determination of PSA in a biological sample. In the assay, the test sample is combined with a labelled conjugate and the response of the conjugate is compared to a standard curve established by performing the assay on one or more calibrators using rPSA (or an immunologically reactive fragment).
                                                                      Human; prostate pharmacogenomic
                                                                                                                                             ABV07911 standard; cDNA; 171 BP
                                                                                                                                                                                                                                                  Sequence 93
                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                                                                                                                                                                                                                                             Recombinant fragments -
                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                    Barnett
                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-APR-1995;
03-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP725139-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Forward primer F2
        20-FEB-2001; 2001WO-US05171.
                        23-AUG-2001.
                                       WO200160860-A2
                                                                                                              13-SEP-2002
                                                                                                                              ABV07911;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-JAN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         baculovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                   (FARB ) BAYER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymerase chain
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                                                                                                                                                                                     16
                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                   μ,
                                                                                                                                                                                                                                                                                                                                                                                                     1996-356106/36
                                                                                                                                                                                                 ATGTGGGTCCCGGTTGTCTTCCTC 24
                                                                                              prostate
                                                                                                                                                                                                                                                                                                                                                     3; Page 5; 19pp; English.
                                                                                                                                                                                                                                                 BP; 12 A; 23 C;
                                                                                                                                                                                                                                                                                                                                                                            prostate-specific antigen (PSA) and immunological for use as an immunoassay calibrator in commercia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transfer
                                                                                                                                                                                                                   Conservative
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                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                             expression
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95US-0383111
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                                                                      marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                   PC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for prostate specific antigen.
                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vector;
                                                                                                                                                                                                                                                                                                                                                                                                                   Yeung
                                                                                              marker cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; amplify
r; PCR; PSA; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                    줐
                                                                                                                                                                                                                                                 31 G; 27 T; 0 other;
                                                                                                                                                                                                                   0
                                                                                                                                                                                     39
                                                                                                                                                                                                                          Score 24; Pred. No. (
                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amplify; prostate specific antigen;
                                                                              carcinogen; pharmacodyanamic
                                                                                              7902
                                                                                                                                                                                                                          0.17;
                                                                                                                                                                                                                                 DB 17;
                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                             commercial
                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                             PSA
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                                                                              marker,
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ABV07396
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Best Local S
Matches 24
 16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                                  17-FEB-2000;
                                                                                                                                                                                                                                             Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   patient; (I) is a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a nucleotide sequence given in Te
specification or its complement.
(a) assessing whether a patient i
                                                                                                                20-FEB-2001;
                                                                                                                                                                              WO200160860-A2
                                                                                                                                                                                                                                                                                            Human prostate
                                                                                                                                                                                                                                                                                                                            13-SEP-2002
                                                                                                                                                                                                                                                                                                                                                            ABV07396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, uses
                                                                                                                                                23-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                            ABV07396 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (e) selecting a composition for inhibiting prostate cancer in a patient;(f) assessing the prostate cell carcinogenic potential of a compound;(g) determining whether prostate cancer has metastasized in a patient;(h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-662795/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (a) assessing whether a patient is afflicted with prostate cancer;(b) monitoring the progression of prostate cancer in a patient;(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 ATGTGGGTCCCGGTTGTCTTCCTC
                                                                                                                                                                                                              sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 1267; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽,
2000US-183319P.
2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-2119007P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-219007P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP;
                                                                                                                 2001WO-US05171
                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        presence of prostate cancer, stage of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             relates
                                                                                                                                                                                                                                                                               expression marker cDNA 7387.
                                                                                                                                                                                                                                marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                            CDNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     55 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     51
                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G; 43 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (I) is useful for:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid molecule (I) comprising 1-9 (ABY00010-ABY62213) of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 other;
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                                                                                                                                                                                                                                                             pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 171;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tate of
useful
cer -
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RESULT 11
ABV373300
ID ABV37
XX
AC ABV37
AC ABV37
XX 16-SE
XX Human
XW Human
XW Homo
XX Homo
OS Homo
OS Homo
OS 23-At
XX 20-FE
PR 11-FE
PR 11-FE
PR 11-FE
PR 11-W
PR 19-JU
PR 18-JU
PR 
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Best Local
                                                                                                                                                                   17-FEB-2000; 2000US-183319P.
16-MAR-2000; 2000US-189865P.
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
13-DEC-2000; 2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV02213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 308 BP; 65 A; 100 C; 79 G; 63 T; 1 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (g) determining whether prostate cancer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 1194; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, used for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Schlegel R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human prostate expression marker cDNA 37321.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABV37330 standard; cDNA; 428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    \left(d\right) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient:
                                                         Schlegel R,
                                                                                                                                                                                                                                                                                                                                                                                                                            23-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200160860-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; prostate cancer; cytostatic; pharmacogenomic marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                     20-FEB-2001; 2001WO-US05171
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24; Conserv
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                                                                                                               MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Endege WO,
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                                                            Monahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        carcinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     . 0.19; 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pharmacodyanamic marker;
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The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, used for detecting presence of prostate cancer, stage of prostate cancer
                                                                                                                                                                                                                                                                                                                                                                       (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient; in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 7668; 11750pp; English.
                                                                                                                                   Sequence 428 BP; 82 A; 132 C; 123 G; 91 T; 0 other;
                                                                                                                                                                                                                                                 (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (g) determining whether prostate cancer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a

    is also useful as a pharmacodyanamic or pharmacogenomic marker.

                                            Similarity
             100.0%; ilarity 100.0%; Conservative 0;
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Pred. No.
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Matches Query Match Local 294

밁 Qy

RESULT 12 AAA61482 23-OCT-2000 AAA61482 standard; DNA; 517 Prostate specific antigen gene fragment. (first entry) ВÞ

PSA gene al DNA amplification; template; prostate sp specific antigen;

nucleic acid

synthesis; human;

detection;

Homo sapiens.

18-MAY-2000

WO200028082-A1.

09-NOV-1998; 08-NOV-1999; 98JP-0317476

99WO-JP06213

(EIKE) EIKEN KAGAKU KK

Notomi T, Hase T;

WPI; 2000-376587/32

Nucleic acid synthesis using primer containing a sequence identical to one on the template for isothermal amplification using simple reagents

Example 8; Fig 17; 95pp; Japanese.

CCCCXXXTT PTT PTXXX PXXX PTXXX The invention relates to a novel method of isothermal DNA amplification. A region is produced at the 3' end of a target sequence which contains sub-sequences designated F1, F2C and F1C, where F1C is the complement of F1. A loop is formed by annealing of F1 to its complementary sequence

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RESULT 13
AAH41845
The resent invention describes a new method for synthesising nucleic acids comprising: (a) preparing a nucleic acid, with an F2c region, able to form a loop by annealing the F1 region of the 3' end to the F1c region on the same chain; (b) synthesis of a complementary chain using the 3' end of F1 as the starting point; (c) replacing the complementary chain by annealing an oligonucleotide containing F2, (complementary to F2c) at its 3' end and using it as a starting point for complementary chain synthesis by a polymerase; and (d) replacing the chain synthesised in step c by annealing a polynucleotide containing a region complementary to a part of the chain and using it 3' end as the starting point. Also described are: (1) a kit for synthesising nucleic scacids; and (2) a kit for detecting target base sequences. The present sequence represents a prostate specific antigen related nucleotide sequence, which is used in an example from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                           Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthesizing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EIKE ) EIKEN KAGAKU KK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prostate specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prostate specific antigen related nucleotide sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 517
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                                                                                                                                                                                                                                                                                                                                                                                                                  8; Fig 17; 96pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antigen; nucleic acid
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Pred. No. 0.2
); Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             synthesis; detection;
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Sequence 517

BP;

94 A; 163 C;

151 G;

109 T; 0 other;

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ATGTGGGTCCCGGTTGTCTTCCTC

Matches

24;

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RESULT 14
ABV25078
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Matches 24
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Best Local
                                                                                                                                                                    The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                         16-MAR-2000;
25-MAY-2000;
                                                                                                                                       cancer in a patient; cancer in a patient;
                                                                                                                                                                                                                                                               Novel isolated nucleic acid molecule associated with cancerous prostate cells and correlating with presence of prostate cancer for detecting presence of prostate cancer, stage of prostate cancer.
                                                Sequence
                                                                                                                                                                                                                                                    Claim 1; Page 4874; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                               Schlegel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-SEP-2002
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                                                                                patient;
                                                                                                               (e) selecting
(f) assessing
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18-JUL-2000;
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                                                                                                                                                                                                                                                                                                                         2001-662795/76
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                                                                    also useful as a pharmacodyanamic or pharmacogenomic
               Similarity
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                                                841 BP;
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2000US-211314P.
2000US-219007P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                               Endege WO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US05171
                                                                                       a composition for inhibiting prostate cancer the prostate cell carcinogenic potential of a ng whether prostate cancer has metastasized in the aggressiveness or indolence of prostate c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              marker;
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                                                                                                                                                                                                                                                                                                                                                                     PREDICTIVE
              100.0%;
                                             A; 262 C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic;
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Score 24; DB 23;
Pred. No. 0.21;
Pred. no. 0.31;
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Pred. No. 0
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                                             262 G; 173 T; 0 other;
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ce of prostate cancer in a
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В

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us-09-829-004a-7.rng
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Search completed: November 19, 2002, 03:06:36 Job time: 171.462 secs
                                                                                                                                                 B
                                                                                     Query Match 100.0%; Score 24; DB 21; Length 990; Best Local Similarity 100.0%; Pred. No. 0:21; Matches 24; Conservative 0; Mismatches 0; Indels
                                                                                                                                                    The invention relates to a method of immunization against prostatic cancer (PC) by administering a nucleic acid (this sequence) that encodes prostate-specific antigen (PSA), or its immunologically active fragment, cells and then expressed to generate an immune response against PSA. The method is used to protect against, or treat, PC, or more generally to induce a cellular immune response to PSA.
                                                                                                                                  Sequence 990 BP; 209 A; 291 C; 273 G; 217 T; 0 other;
                                                                                                                                                                                                                                                           Disclosure; Page 42; 47pp; English.
                                                                                                                                                                                                                                                                           Immunization against prostatic cancer using nucleic acid encoding prostate-specific antigen, for treatment or prophylaxis -
                                                                                                                                                                                                                                                                                                                       WPI; 2000-072551/06.
P-PSDB; AAY56048.
                                                                                                                                                                                                                                                                                                                                                         Weiner DB, Kim JJ;
                                                                                                                                                                                                                                                                                                                                                                                (UYPE-) UNIV PENNSYLVANIA.
                                            29-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                28-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-DEC-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W09961068-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunization; prostate cancer; prostate-specific antigen; psa; cytostatic; immune response; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human prostate-specific antigen gene.
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a a

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Sequence:
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Gapop 10.0 , Gapext 1.0
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24
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Copyright (c) 1993 - 2002 Compugen Ltd.
           em_gss_mus:*
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Pred. No. and is greater derived is the number of results predicted by chance to have a ater than or equal to the score of the result being printed, rived by analysis of the total score distribution.

SUMMARIES

100.0 408 12 BF678836	100.0 336 9 AA506459 7 7 100.0 371 9 AA503943 7 100.0 391 9 AA506939 7 7 100.0 408 12 BF678836	316 9 AA593245	Result Query No. Score Match Length DB ID Des
BF678836 602153143		AA593245 nn07e10.s	Description

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Q94942	F6810	F67801	F67336	F67380	F96522	F67533	99	F67324	F67532	F67376	F68097	5573	F67807	F67755	F68150	7514	F674	F67551	F67577	F67838	F67574	F96525	F67601	E97398	7916	5472	473	F964	524893	F67760	F679	F67959	E84053	84070	63990	4	E8405	I04647	
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RESULT 1
AA593245
LOCUS
DEFINITION

AA593245 316 bp mRNA linear EST 25-SEP-1997 nn07e10.sl NCI_CGAP_Pr4.1 Homo sapiens cDNA clone IMAGE:1077066 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);

ALIGNMENTS

SOURCE ORGANISM ACCESSION VERSION KEYWORDS TST

human

AA593245.1 GI:2409007

AA593245 mRNA sequence.

REFERENCE AUTHORS TITLE Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 316)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),

COMMENT JOURNAL

National
Tumor Gene Index
AL Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B: Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bio.llnl.gov/bbrp/image/image.html

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REFERENCE
AUTHORS
TITLE
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VERSION
KEYWORDS
SOURCE
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AA506459
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

El (bases 1 to 336)

I (bases 1 to 336)

RNCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

RNCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

Rational Cancer Institute, Cancer Genome Anatomy Project (CGAP),
Tumor Gene Index

Unpublished (1997)

Contact: Robert Strausberg, Ph.D.

Emmall: cgapbs-remail.nih.gov

Tissue Procurement: bavid G. Bostwick, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.

DNA Library Preparation: David B. Krizman, Ph.D.

DNA Library Preparation: David B. Krizman, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/Dbrp/image/image.html
Insert Length: 442

Std Ezror: 0.00

Trocation Constitution information can be constitution information can be constitution information can be constitution.
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High quality sequence stop: 293.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA506459 mRNA linear EST 20-AUG-1997 nh46a08.sl NCI_CGAP_Pr5 Homo sapiens cDNA clone IMAGE:955382 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA506459.1 GI:2242699
EST.
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AA506459
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/lab_host="Organ: prostate; Vector: pAMP10; mRNA made from /note="Organ: prostate; Vector: pAMP10; mRNA made from /note="Organ: prostate intraepithelial neoplasia (high-grade); cDNA made by oligo-dT priming. Non-directionally cloned. Size-selected on agarose gel, average insert size 600 bp. cDNA Library Preparation: David B. Krizman, Ph.D. Reference: Krizman et al. (1996) Cancer Research 56:5380-5383. cDNA Library Arrayed by: Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /tissue_type="prostatic intraepithelial neoplasia - high
grade"
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/sex="male"
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/db_xref="taxon:9606"
/clone="IMAGE:1077066"
                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:955382"
/clone_lib="NCI_CGAP_Pr5"
                        /tissue_type="prostate"
/lab_host="DH10B"
                                                                                                                                                                                  Location/Qualifiers
/note="Vector: pAMP10; mRNA made from normal prostatic
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Дb QΥ

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1 ATGTGGGTCCCGGTTGTCTTCCTC 24
                   1 ATGTGGGTCCCGGTTGTCTTCCTC 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AL Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: Gapbs-remail.nih.gov
Tissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.
CDNA Library Preparation: David B. Krizman, Ph.D.
CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
www-bb.llnl.gov/bbrp/inage/inage/inage/inage
Insert Length: 454 Std Error: 0.00
Seq primer: -40m.3 fwd. ET from Amersham
High quality sequence stop: 365.
High quality sequence stop: 365.
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ATGTGGGTCCCGGTTGTCTTCCTC 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AA503943 371 bp mRNA linear EST 20-AUG-1997 nh38405.s1 NCI_CGAP_Pr5 Homo sapiens cDNA clone IMAGE:954633 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);, mRNA sequence.

AA503943 AA503943 AA503943
                                                                                          l Similarity
24; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 371)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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24; Conservative
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                                                                                       100.0%;
ilarity 100.0%;
Conservative (
                                                                                                                                                                                                                      63
                                                                                                                                                                                                                                                                                                                                                                                     Organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:954633"
/clone="tib="NCI_CGAP_Pr5"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           epithelial cells, cDNA made by oligo-dT priming. Non-directionally cloned. Size-selected on agar average insert size 600 bp. "
a 106 c 99 g 74 t
                                                                                                                                                                                                           /tissue_type="prostate"
//lab_host="DH10B"
/lab_host="DH10B"
/lab_host="DH10B"
/note="vector: pawn10; mRNA made from normal prostatic epithelial cells, cDNA made by oligo-dT priming.

epithelial cells, cDNA made Size-selected on agarose gel, Non-directionally cloned. Size-selected on agarose gel, average insert size 600 bp.

a verage insert size 600 bp.

a verage insert size 600 bp.

b t
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                                                                                                0; Mismatches
                                                                                                                   Score 24;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 24; DB 9; Length 336; Pred. No. 1.9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                         ; DB 9;
. 1.9;
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                                                                                                  0;
                                                                                                                                                Length 371
                                                                                                  Indels
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REFERENCE
AUTHORS
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BF678836
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ORIGIN
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Best Local (
                                                                                                                   TITLE
                                                                                                                                                                                                                   ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          source
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D.

Email: cgapbs r@mail.nih.gov

Tissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: David B. Krizman, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 488 Std Error: 0.00

Seq primer: -40m13 fwd. ET from Amersham.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA506939 391 bp mRNA linear EST 20-AUG-1997 nh64b07.sl NCI_CGAP_Pr8 Homo sapiens cDNA clone IMAGE:957109 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                                                                                                             1 (bases 1 to 408)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                               BF678836
                                                                                                                                                                                                                                                                                                                  mRNA sequence.
                                                                                                                                                                                                                                                                                                                                   602153143F1 NIH_MGC_83 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 391)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA sequence.
                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                         BF678836.1 GI:11952731
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Score 24; Dilarity 100.0%; Pred. No. 1.: Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note-"Vector: pamP10; mRNA made from invasive prostate tumor, cDNA made by oligo-dT priming. Non-directionally cloned. Size-selected on agarose gel, average insert size 600 bp. "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="NCI_CGAP_Pr8"
/sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:957109"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /tissue_type="prostate"
/lab_host="DH10B"
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1.9;
                                                                                                                                                                                                                                                                                                                                 mRNA linear EST 21-DEC-2000 s cDNA clone IMAGE:4294258 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                 Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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AUTHORS
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AA528287
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ORIGIN
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ORGANISM
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Query Match
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                                                                                                                                                                                                                                                                                                                                                              Tunor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
, Michael Emmert-Buck, M.D., Ph.D.
                                                                                                                                                  cDNA Library Preparation: David. 8. Krizman, Ph.D. cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html Seq primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA528287

A57 bp __mrNA linear EST 20-AUG-1997
nh26d08.sl NCI_CGAP_Pr3 Homo sapiens cDNA clone IMAGE:953487
similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 457)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mRNA sequence.
                                                                                                                 High quality sequence stop: 187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA528287.1 GI:2270356
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Plate: LLCM1143 row: e column: 11
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DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab_host="DH10B (TI phage-resistant)"
/note="Organ: prostate; Vector: pDNR-LIB (Clontech);
/note="Organ: prostate; Vector: pDNR-LIB (Clontech);
Site_1: SfiI (ggccgctcggcc); Site_2: SfiI (ggccattatggcc);
5' and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor
sequence: 5'-ATTCTACAGGCCAGCGGCCACATG-dT(30)BN-3'
(where B = A, C, or G and N = A, C, G, or T). Average
insert size 1.4 kb (range 0.5-4.0 kb). 14/15 colonies
contained inserts by PCR. This library was enriched for
full-length clones and was constructed by Clontech
Laboratories (Palo Alto, CA)."

74 a 127 c 115 g 92 t
/organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
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/clone="IMAGE:4294258"
/clone_lib="NIH_MGC_83"
                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ocation/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
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/clone="IMAGE:953487"

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1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 465)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Magai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and Simpson, D. H., Sares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and Simpson, D. H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      465 bp mRNA linear
MR3-FN0209-070201-010-b11 FN0209 Homo sapiens cDNA,
BI046475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010,
                                                                                                                                                                                                                                                                                                                                                                                 Dimail: asimpson@ludwig.org_br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR3&t2=MR3-FN0209-
070201-010-bilstj=2001-02-07&t4=1)
Seq_primer: puc 18 forward
seq_primer: puc 18 forward
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+55-11-2707001
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Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Natl. Acad. Sci. U.S.A.
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/lab_host="DH10B"
   /note="Organ: prostate_normal; Vector: pucl8; Site_1: SmaI; site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse
Research) profiles into
                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="FN0209"
/dev_stage="Adult"
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Pred. No. 1.9;
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RESULT 7
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LOCUS
DEFINITION

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26

ACCESSION VERSION KEYWORDS

SOURCE ORGANISM

REFERENCE

COMMENT

source

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JOURNAL MEDLINE

BASE COUNT ORIGIN

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KEYWORDS
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Best Local S
Matches 24
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1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutherla; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 495)
Dias Neto, E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva-W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher;P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scritpts/gethtml2.pl?tl=&t2=RC1-FN0188-260
700-022-f06xt3-2000-07-5&st4-=1)
Seq primer: puc 18 forward

Seq primer: puc 18 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: +55-11-2704922
Fax: +55-11-2707001
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Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Simpson A.J.G.
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                                                                                                                                  /note="Organ: prostate_normal; vector: puc18; Site_1: Smal | Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Pattent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
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/db_xref="taxon:9606"
/clone_lib="FN0188"
/dev_stage="Adult"
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1 (bases 1 to 505)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
AA639901.1 GI:2563680
EST.
                                                                             np08f03.s1 NCI_CGAP_Pr3 Homo s
similar to gb:M21895 PROSTATE
                                      mRNA sequence.
AA639901
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                      AA639901
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700-021-a02&t3=2000-07-21&t4=1)
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n quality sequence stop: 505.
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                                                                                                                                                                                                                                                                                                                                                                                         /note-"Organ: prostate_normal; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
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/clone_lib="FN0188"
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Pred. No.
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                                                                   שע Linear EST 23-OCT-1997 sapiens cDNA clone IMAGE:1115741 SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                                                                                                                                                                                                                                                                                                                  DB 12;
1.9;
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                                                                                   Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,Brunstein,A., deollveira,P.S., Bucher,P., Jongeneel,C.V., O'Ha,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 552)

Dias Neto,E., García Correa,R., Verjovski-Almeida,S., Briones,M.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tissue Procurement: W. Marston Linehan, M.D., Rodr
, Michael Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David, B. Krizman, Ph.D.
cDNA Library Arrayed by: Genome Systems Inc., Gre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                        Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                                                                                                                                                                                                                                            EST
                                                                                                                                                                                                                                                                                                                                                                                          BE840701 552 bp mRNA linear RC1-FN0188-140800-011-g05 FN0188 Homo sapiens cDNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia; Eutheria; Primates;
1 (bases 1 to 523)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Seq primer: -40ml3 fwd. ET from Amersham
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    National Cancer Institute, Cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note="Vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st strand cDNA was primed with oligo(dT),17 on 50 ng of DNAse-treated, total cellular RNA obtained from 5,000-10,000 microdissected cells histologically-determined to be fully malignant prostate cancer cells. Double-stranded cDNA was ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
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/lab_host="DH10B"
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/db_xref="taxon:9606"
/clone="IMAGE:1115741"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="NCI_CGAP_Pr3"
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Acad. Sci. U.S.A.
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Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagal N.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Roldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BE840537
RC1-FN0188-260700-022-h06
BE840537
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This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=st2=RCl-FN0188-140
800-011-g05st3=2000-08-14st4-1)
seq primer: puc 18 forward
High quality sequence start: 2
High quality sequence start: 2
Location/Qualifiers
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Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
Fax: +55-11-2707001
Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
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                                                                                                                                           Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 an
                                                                                                                                                                                                                     Contact: Simpson A.J.G.
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                                                                                             +55-11-2704922
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//db_xref="taxon:9666"
//db_xref="taxon:9666"
//db_xref="taxon:9666"
//dev_stage="Adult"
//dev_stage="Adult"
//note="Organ: prostate_normal; Vector: puc18; Site_1: SmaI; A mini-library was made by cloning a products derived from ORESTES PCR (U.S. Lietters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."
79 a 150 c 171 g 152 t
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FN0188 Homo sapiens
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Tissue Procurement: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLCM1145 row: a column: 10
High quality sequence stop: 587.
Location/Qualifiers
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1 (bases 1 to 591)

NIH-MGC http://mgc.nci.nih.gov/...
National Institutes of Health, Mammalian Gene Collection (MGC) Unpublished (1999)

Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BF679591 591 bp mRNA linear EST 21-D 602154033F1 NIH_MGC_83 Homo sapiens cDNA clone IMAGE:4294929
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High quality sequence stop: 572
Location/Qualifiers
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/clone=lib="NIH_MGC_83"
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/site_1: SfiI (ggccgcttatggcc);
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/site_1: SfiI (ggccgcttatggcc);
/site_1: SfiI (ggccgttatggcc);
/site_1: SfiI (ggccgtatggcc);
/site_1: SfiI (ggccgtatggccgcatggccgcatggccgcatggccgcatggccgcatggccgcatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggccatggcc
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Pred. No. 1.9;
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RESULT 15
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      DEFINITION
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Best Local
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                                                                                                                                                                                      ATGTGGGTCCCGGTTGTCTTCCTC 66
BF677605 648 bp r 602085679F1 NIH_MGC_83 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 638)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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EST.
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BF679511
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602153922F1 NIH_MGC_83 Homo sapiens
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ilarity 100.0%;
Conservative (
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/lab_host="DH10B (TI phage-resistant)"
/note="Organ: prostate; Vector: pDNR-LIB (Clontech);
/note="Organ: phage-resistant;"
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a 181 c 166 g 12
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/db_xref="taxon:9606"
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   mRNA linear EST 21-DEC-2000 cDNA clone IMAGE:4249939 5',
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Search completed: November 19, 2002, 05:17:29 Job time : $1377.08 \ \text{secs}$

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REFERENCE
AUTHORS
TITLE
JOURNAL
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KEYWORDS
SOURCE
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                                                                                                               Query Match
Best Local Similarity
                                                                                         Matches
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43 ATGTGGGTCCCGGTTGTCTTCCTC
                        1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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Tissue Procurement: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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BF677605
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Robert Strausberg, Ph.D:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
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                                                                                                                                                                                                              124 a
                                                                                         Conservative
                                                                                                                                                                                                                                                   /note="organ: prostate; Vector: pDNR'LIB (Clontech); Site_1: Sfil (ggccqctcggcc); Site_2: Sfil (ggccqttatggcc); Site_2: Sfil (ggccqttatggcc); S' and 3' adaptors were used in cloning as follows: 5' adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGACGGCGACATG-dT(30)BN-3' (where B = A, C, or G and N = A, C, G, or T). Average insert size 1.4 kb (range 0.5-4.0 kb): 14/15 colonies contained inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech
                                                                                                                                                                                                         Laboratories (Palo Alto, CA)."
198 c 185 g 141 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /clone="IMAGE:4249939"
/clone_lib="NIH_MGC_83"
/lab_host="DH10B (T1 phage-resistant)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                               100.0%;
                                                                                         0;
                                                                                                               Score 24; D
Pred. No. 2;
                                                                                         Mismatches
                                                                                                                                        DB
                                                                                                                                     12;
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                                                                                                                                     Length 648;
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                                                                                         Gaps
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Title:
Perfect score:
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         Issued_patents_NA:*
1: /cgn2_6/ptodata/2/
2: /cgn2_6/ptodata/2/
3: /cgn2_6/ptodata/2/
4: /cgn2_6/ptodata/2/
5: /cgn2_6/ptodata/2/
6: /cgn2_6/ptodata/2/
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length: 2000000000
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Match
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1 atgtgggtcccggt
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156:345 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gapop 10.0 , Gapext 1.0
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Listing first 45 summaries
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
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/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/2/ina/backfiles1.seq:*
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US-08-454-720A-42
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US-08-605-785-607-
US-08-605-785-607-
US-08-96-77-492-31
US-09-079-630-31
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US-09-079-630-31
US-09-079-630-31
US-08-454-720A-38
US-08-68-982-075D-10
US-08-768-859A-5
US-08-767-820A-5
US-08-790-137-2
US-09-102-732-2
US-09-261-767-2
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		Sequence 20, Appl	Sequence 20, Appl	Sequence 20, Appl	-	-		Sequence 2, Appli	•	•	Sequence 99, Appl	Sequence 1, Appli	Sequence 4, Appli	Sequence 7, Appli	Sequence 8, Appli	Sequence 6, Appli	Sequence 5, Appli	Sequence 7, Appli

Minimum Maximum

DB DB

Database

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Sequence 42, Application US/08454720A
Patent No. 5766888
GENERAL INFORMATION:
APPLICANT: Sobol, Robert E.
APPLICANT: Green, Mark R.
APPLICANT: Kawasaki, Ernest S.
TITLE OF INVENTION: Detection of Car
TITLE OF INVENTION: Nucleic Acid Amm
NUMBER OF SEQUENCES: 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
OTHER INFORMATION: Description of Artificial
TOTHER INFORMATION: oligonucleotide
US-09-210-016-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 70
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/09210016 Patent No. 6221579 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Everhart, Dennis S.
APPLICANT: Kaylor, Rosann M.
APPLICANT: McGrath, Kevin
TITLE OF INVENTION: Patterned Binding of Functionalized Microspheres
TITLE OF INVENTION: Optical Diffraction-Based Biosensors
FILE REFERENCE: 11301-0880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/210,016; AND CURRENT FILING DATE: 1998-12-11
NUMBER OF SEQ ID NOS: 2
                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Hoffman-La
STREET: 340 Kings
CITY: Nutley
STATE: New Jersey
                     ADDRESSEE: HOLLINGSLAND
                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                     1 ATGTGGGTCCCGGTTGTCTTCCTC
                                                                                                                                                                                                                                                                                                                      ATGTGGGTCCCGGTTGTCTTCCTC
                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                      Roche Inc.
Street
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Pred. No: 0
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ZIP: 07110
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy

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HOLECULE TYPE: DNA (genomic)
US-08-454-720A-42
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INFORMATION FOR SEQ ID NO: 42
SEQUENCE CHARACTERISTICS:
LENGTH: 111 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 24; Best Local Similarity 100.0%; Pred. No. Matches 24; Conservative 0; Mismatch
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-LOS/MS-DOS
SOFTWARE: Patentin Release #1.01, version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/454,720A
FILING DATE: May 31, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNDER: 08/357,565
APPLICATION UNDER: 08/357,565
APPLICATION UNDER: 08/357,565
APPLICATION UNDER: 35321
REGISTRATION UNDER: 35321
REGISTRATION UNDER: 35321
REGISTRATION UNDER: 35321
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Patent No.
APPLICATION NUMBER: US/08/358
FILING DATE: 15-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION UNMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-
TELEOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3439
TELEPHONE: 215-568-3439
                                                                                                                                                                                                                    STATE: PA
ZIP: 19103
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-TOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: MULHOLland, S. Grant
APPLICANT: Moreno, Jose
APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis of
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz &
STREET: One Liberty Place 46th. Floor
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 91
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 814-2974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NERAL INFORMATION:
APPLICANT: Croce, Carlo
APPLICANT: Gomella, Leor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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9178
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.023;
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1 ATGTGGGTCCCGGTTGTCTTCCTC 24

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; MOLECULE TYPE: DNA (
; HYPOTHETICAL: NO
; ANTI-SENSE; NO
US-08-764-527A-13
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; MOLECULE TYPE: DNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-08-358-782D-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 4
US-08-764-527A-13
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    Query Match
Best Local S
Matches 24
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Patent No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM.
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATES: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 992 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Croce, Carlo
APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. Grant
APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Wethods of Detecting Micrometastasis of
TITLE OF INVENTION: Prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CILL PA
STATE: PA
STATE: 19103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & STREET: One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION:
h 100.0%; Score 24; DB 2; Similarity 100.0%; Pred. No. 0.028; 24; Conservative 0; Mismatches
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es ": 0; Indels
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                                                                            Length 992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                      tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 1462 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
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                                            FITLE OF INVENTION:
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                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: One Landelphia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                            1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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                                                                                                                                                    INFORMATION:
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24; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beardell, Lori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/08358782D
                                                                                                                                                                                    Application US/08764527A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Woodcock Washburn Kurtz Mackiewicz & One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          215-568-3439
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                                                                           Fischer, Rainer
                                                                                                      Mulholland, S. Grant
                                                                                                                       Gomella, Leonard
                                                                                                                                       Croce, Carlo
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                                                                                       Moreno, Jose
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SYSTEM: PC-DOS/MS-DOS
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Woodcock Washburn Kurtz Mackiewicz &
                                                                                                                                                                                                                                                                                                                                                                                                                   NO
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N: 435
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                                                           Methods of Detecting Micrometastasis
                                            Prostate
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Pred. No. 0
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                                                                                                            SEQ ID NO 31
LENGTH: 7130
TYPE: DNA
ORGANISM: Homo sapiens
                                   Matches
                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                         GENERAL
                                                                                                                                                                                                                                                                                                                                                                      Sequence 31, Appli
Patent No. 6287569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/056,105 CURRENT FILING DATE: 1998-04-06
                                                                                                                                                                                                                             CURRENT FILING DATE: 1998-04-06 EARLIER APPLICATION NUMBER: 60/043,467
                                                                                                                                                                                                                                                                            FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                       APPLICANT: KIPPS, THO APPLICANT: WU, YUNQI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 215-568-3439 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                             EARLIER FILING DATE: 1997-04-10 NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: VACCINES WITH ENHANCED INTRACELLULAR TITLE OF INVENTION: PROCESSING
                                                                                                                                                                                 SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 15-DEC-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25 ATGTGGGTCCCGGTTGTCTTCCTC
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REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/764,527A
   1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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                                   l Similarity
24; Conserv
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                                                                                                                                                                                FastSEQ for Windows Version 3.0
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                                   Conservative
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15-DEC-1994
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Pred. No. 0.028;
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                                 Mismatches
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                                                  .032;
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US-08-394-033-1
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CURRENT FILING DATE: 2000-06-27
NUMBER OF SEQ ID NOS: 835
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 607
LENGTH: 22
Typer: ....
                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/08394033
Patent No. 5614372
GENERAL INFORMATION:
APPLICANT: Lilja, Hans
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Best Local 9
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                   APPLICANT: Lilja, Hans
APPLICANT: Lundwall, Ake
APPLICANT: Lundwall, Ake
APPLICANT: Lowgren, Janita
TITLE OF INVENTION: Early Detection of Prostate Cancer (CAP)
TITLE OF INVENTION: by Employing Prostate Specific Antigen (PSA) and Human
TITLE OF INVENTION: candular Kallikrein (HGK-1)
                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue, NW, Suite 1000
CURRENT APPLICATION DATA APPLICATION NUMBER: U
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                                                                                                                                                               CITY: Washington
STATE: DC
                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ATGTGGGTCCCGGTTGTCTT 20
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20; Conserv
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Carter, Darrick
Li, Samuel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fanger, Gary R. Retter, Marc W. Stolk, John A.
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Kalos, Michael
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                                                                                                                                            USA
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ick, Thomas S
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100.0%; Pr
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US/08/394,033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 20; DB 4; Length 22; pred. No. 1.3; 0; Mismatches 0; Indels
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; ORIGINAL SOURCE:
; ORGANISM: Home
US-08-394-033-1
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US-08-577-492-31
ATTORNEY/AGENT INFORMATION:
NAME: Cherry, David A.
REGISTRATION NUMBER: 35,099
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEPAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 31:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARRCTERISTICS:
LENGTH: 28 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Description Descri
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFFWARE: WordDerfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/577,492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                        FILING DATE: 22-DEC-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA: APPLICATION NUMBER: GB 9426227.6
FILING DATE: 23-DEC-1994
PRIOR APPLICATION DATA: 9512996.1
APPLICATION NUMBER: GB 9512996.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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HYPOTHETICAL: NO
ANTI-SENSE: NO
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NAME: Ihnen, Jeffrey L.
REGISTRATION UNMBER: 28,957
REFERENCE/DOCKET NUMBER: 23
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-962-4810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock, Washburn, Kurtz, Mackiewicz
STREET: One Liberty Place, 46th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Perry, Martin John
APPLICANT: Lumb, Simon Mark
TITLE OF INVERVION: HUMAN PHOSPHODIESTERASE TYPE IVC, AND
TITLE OF INVENTION: ITS PRODUCTION AND USE
NUMBER OF SEQUENCES: 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: UZIP: 19103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: One Liberty
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11 ATGTGGGTCCCGGTTGTC 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 ATGTGGGTCCCGGTTGTC 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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N: 435
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26-JUN-1995
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TYPE: nucleic acid STRANDEDNESS: double

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US-09-079-630-31
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                                                                                                                           US-09-079-630-31
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GENERAL INFORMATION:
                                                             Matches
                                                                                           Query Match
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                                                                               Best
                                                                                                                                                                                                     TELEFAX: (215) 568-3439 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 19103
COMPUTER READABLE FORM:
TYPE: Floppy disk
                                                                                                                                                                                                                                                  NAME: Cherry, David A.
REGISTRATION NUMBER: 35,099
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                          FILING DATE: 23-DEC-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 7
                                                                                                                                                                                                                                                                                                                FILING DATE: 26-JUN-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/577,492
FILING DATE: 22-DEC-1995
APPLICATION NUMBER: GB 9426227.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOCTWARE: Wordperfect 6.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: HUMAN PHOSPHODIESTE TITLE OF INVENTION: ITS PRODUCTION AND NUMBER OF SEQUENCES: 40.

CORRESPONDENCE ADDRESS:
198 GGTCCCGGTTGTCTTCCT 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  198 GGTCCCGGTTGTCTTCCT 215
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                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
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                                                                                                                                                                                       LENGTH:
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                              6 GGTCCCGGTTGTCTTCCT 23
                                                               18;
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Similarity 100.0%;
                                                                            Similarity
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                                                                                                                                                                      nucleic acid
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                                                                                                                                                                                      2153 base pairs
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One Liberty Place, 46th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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Lumb, Simon Mark
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                                                                                                                                          linear
                                                                                                                                                         double
                                                                              100.0%;
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                                                                              Score 18; Pred. No.
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Pred. No
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                                                             Mismatches
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15;
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                                                                                           Length 2153;
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RESULT 12

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                                                                                                                                                                                                                                                                                                                                                            Sequence 38, Application US/08454720A Patent No. 5766888
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 73.3%;
Best Local Similarity 83.3%;
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
COMPUTER: READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/99/042,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                  APPLICANT: Sobol, Robert E.
APPLICANT: Green, Mark R.
APPLICANT: Kawasaki, Ernest S.
TITLE OF INVENTION: Detection of Ca
TITLE OF INVENTION: Nucleic Acid An
NUMBER OF SEQUENCES: 42
CORRESSPONDENCE ADDRESS:
ADDRESSEE: Hoffman La Roche Inc.
                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Sobol,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (206) 682-603
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 1548 base pair:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Burian, Jan
APPLICANT: Kay, William W.
TITLE OF INVENTION: REPLICAN
TITLE OF INVENTION: CONTROLL
TITLE OF INVENTION: CONTROLL
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                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            411 ATGTGGCTTCCGGTTCTCTCCGC 388
             COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                          STATE: New Jersey COUNTRY: U.S.A. ZIP: 07110
                                                                                                                                               STREET: 340 K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
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REFERENCE/DOCKET NUMBER: 66
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FILING DATE: 13-MAR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ATGTGGGTCCCGGTTGTCTTCCTC 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
DEDNESS: single
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                                                                                                                                                                  340 Kingsland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
             IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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PatentIn Release #1.01,
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                                                                                                                                                                                        Roche Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
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Pred. No. 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       701 Fifth Avenue
                                                                                                                                                                                                                                             Amplification
                                                                                                                                                                                                                                                                  Carcinoma Metastases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Version #1.30
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/454,720A
FILING DATE: May 31, 1995
CLASSIFICATION: 435

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RESULT 14

US-08-983-075D-10

Sequence 10, Application US/08983075D

Patent No. 6303361

Patent No. 6303361

GENERAL INFORMATION:

APPLICANT: VIHKO, PIIKKO

ITITE OF INVENTION: HUMAN GLANDULAR KALLIKREIN-1 (HK2)

FILE REFERENCE: 1491/44025

CURRENT APPLICATION NUMBER: US/08/983,075D

CURRENT APPLICATION NUMBER: PCT/F196/00382

PRIOR APPLICATION NUMBER: PCT/F196/00382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: December 16, 1994
ATTORNEY/AGENT INFORMATION:
NAME: PELTY, DOUGlas
REGISTRATION NUMBER: 35321
REFERENCE/DOCKET NUMBER: 9178
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 814-2977
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 1864 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
US-08-454-720A-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; NAME/KEY: misc_feature
; OTHER INFORMATION: nucleotide sequence encoding the prepro-sequence
; OTHER INFORMATION: of the hK2 protein
US-08-983-075D-10
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US-09-385-982-435/c
Sequence 435, Application US/09385982
Patent No. 6262334
GENERAL INFORMATION:
APPLICANT: ENDEGE, WILSON O., ET AL.
APPLICANT: ENDEGE, WILSON O., ET AL.
TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
TITLE OF INVENTION: PRODUCTS: II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 68.3
Best Local Similarity 94.4
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
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CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: 08/357,565
APPLICATION NUMBER: 16, 1994
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Pred. No. 76;
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Pred. No. 89;
0; Mismatches
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CURRENT APPLICATION NUMBER: US/09/385,982
CURRENT FILING DATE: 1999-08-30
EARLIER APPLICATION NUMBER: 09/328,111
EARLIER APPLICATION NUMBER: 60/238,111
EARLIER APPLICATION NUMBER: 60/117,393
EARLIER FILING DATE: 1999-06-08
EARLIER FILING DATE: 1999-01-27
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER FILING DATE: 1999-01-27
EARLIER APPLICATION NUMBER: 60/098,639
EARLIER FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 544
SOFTMARE: FASTSEQ for Windows Version 3.0
INUMBER OF SEQ ID NOS: 544
SOFTMARE: FASTSEQ for Windows Version 3.0
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:/cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
:/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
:/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
:/cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
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:/cgn2_6/ptodata/1/pubpna/US08_NEW_PUB.seq:*
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:/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
:/cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
:/cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
:/cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
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:/cgn2_6/ptodata/1/pubpna/US00_PUBCOMB.seq:*
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Copyright (c) 1993 - 2002 Compugen Ltd.
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110 US-09-755-100-2
110 US-09-759-143-607
110 US-09-780-669-607
110 US-09-784-808-429
110 US-09-974-300-2790
110 US-09-974-301-16219
110 US-09-974-301-16219
110 US-09-974-301-277
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US-09-755-100-4
US-09-755-100-5
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Sequence 7, Appli
Sequence 4, Appli
Sequence 5, Appli
Sequence 3, Appli
Sequence 1, Appli
Sequence 607, App
Sequence 607, App
Sequence 2790, App
Sequence 429, App
Sequence 4, Appli
Sequence 277, Ap
Sequence 2277, Ap
Sequence 3, Appli
Sequence 379, App
Sequence 379, App
Sequence 20503, A
Sequence 13038, A
Sequence 13058, A
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RESULT 1 US-09-829-004A-7

ALIGNMENTS

Sequence 7, Application US/09829004A Patent No. US20020132976A1

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US-09-918-063-31 US-09-918-063-73	-063	US-09-918-063-14	US-09-777-564-1235 US-09-736-457-1047	-564	-09-918	US-10-044-090-347		US-09-925-300-153	US-09-938-642-1	US-09-755-100-6	US-09-974-300-1552	US-09-880-107-2307	-09	US-09-956-999-7	US-09-925-297-286	US-09-960-352-4535	US-10-033-528-335	ő	US-09-960-352-257	US-09-960-352-11130	US-09-960-352-12564
Sequence 31, Appl Sequence 73, Appl	29,	Sequence 14, Appl	Sequence 1235, Ap Sequence 1047, Ap	Sequence 67, Appi Sequence 699, App		347,	348,	153,	<u>,</u>	Sequence 6, Appli	Sequence 1552, Ap	2307,	Sequence 101, App		Sequence 286, App	Sequence 4535, Ap	335,	Sequence 335, App	Sequence 257, App	Sequence 11130, A	Sequence 12564, A

PRIOR APPLICATION NUMBER: US 60/195,456 PRIOR FILING DATE: 2000-04-10 NUMBER OF SEQ ID NOS: 12 SOFTWARE: PATENTIN VERSION 3.0 SEQ ID NO 7 LENGTH: 24 TYPE: DNA ORGANISM: Artificial Sequence FEATURE: RESULT 2 US-09-755-100-4 ₽ Qy ; OTHER INFORMATION: CLP313 US-09-829-004A-7 Sequence 4, Application US/09755100 Patent No. US20020099189A1 GENERAL INFORMATION: APPLICANT: SAVITZKY, Kinneret et al. TITLE OF INVENTION: NOVEL NUCLEIC ACID FILE REFERENCE: 2786-0156P GENERAL INFORMATION: APPLICANT: Chong, Pele APPLICANT: Pedyczak, Artur APPLICANT: 914, Charles Dwo Yuan TITLE OF INVENTION: Immunogenic Peptides Derived TITLE OF INVENTION: Uses Thereof FILE REFERENCE: 11014-24 CURRENT APPLICATION NUMBER: US/09/829,004A CURRENT FILING DATE: 2001-04-16 Query Match 100 Best Local Similarity 100 Matches 24; Conservative 100.0%; Score 24; DB 10; Length 24; 100.0%; Pred. No. 0.016; 0; red. No. 0.(AND AMINO ACID SEQUENCES from Indels Prostate-Specific 0 Gaps Antigen 0 P

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PRIOR APPLICATION NUMBER: IL 12587
PRIOR APPLICATION NUMBER: IL 12587
PRIOR PRILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 12
SOCTWARE: Patentin Ver. 2.1
SEQ ID NO 5
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CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: IL 128587
PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
                             US-09-755-100-3
Sequence 3. Application US/09755100
Sequence 3. Application US/09755100
Patent No. US20020099189A1
GENERAL INFORMATION:
APPLICANT: SAUTIZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156p
CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT PAPLICATION SUMBER: US/09/755,100
PRIOR APPLICATION NUMBER: IL 128587
PRIOR APPLICATION NUMBER: 1599-02-18
PRIOR APPLICATION NUMBER: 1120439
PRIOR APPLICATION NUMBER: 1120439
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; ORGANISM: Homo sapiens
US-09-755-100-5
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Patent No. US20020099189A1
GENERAL INFORMATION:
APPLICANT: SAVITZRY, Kinneret et al.
APPLICANT: SAVITZRY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION UNMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
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ORGANISM: Homo sapiens
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IL 131363
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0.028;
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; TYPE: DNA; ORGANISM: Homo sapiens US-09-755-100-1
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US-09-755-100-2
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US-09-755-100-1
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Patent No. US20020099189A1
GENERAL INFORMATION:
APPLICANT: SAVITZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
CURRENT FILING DATE: 2001-01-08
                                                                                                                                                                                                                                                      Sequence 2, Application US/09755100

Patent No. US20020099189A1

GENERAL INFORMATION:
APPLICANT: SAVITZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
CURRENT FILING DATE: 2001-01-08
                                                     PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PATENTIN VET. 2.1
SEQ ID NO 2
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SOFTWARE: PatentIn Ver. 2.1
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NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
LENGTH: 3846
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PRIOR FILING DATE: 1999-02-18
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PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
PRIOR FILING DATE: 1999-08-11
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PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-04-14
ORGANISM: Homo sapiens
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                                         ENGTH: 4661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100 hes 24; Conservative
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OS: 12
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Pred. No. 0.029;
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RESULT 8
US-09-780-669-607
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SEQ ID NO 607
                                                                                                                                          Patent No.
                                                                                                                                                     Sequence 607,
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Patent No. US20020022248A1
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                                       APPLICANT:
                                                               APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.427C23
CURRENT FILING DATE: 2001-01-12
CURRENT FILING DATE: 2001-01-12
CURRENT OF SEQ ID NOS: 934
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APPLICANT:
APPLICANT:
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TYPE: DNA
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                                                                                                                                INFORMATION:
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                                                                                                                                         07, Application US/09780669
US20020051977A1
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Fanger, Gary R. Retter, Marc W. Stolk, John A.
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Hepler, William
                                                                        Dillon, Davin C.
Mitcham, Jennifer L.
Harlocker, Susan L.
                                     Henderson, Robert A. Kalos, Michael D.
                                                              Jiang, Yuqu
                                                                                                                 Xu, Jiangchun
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Retter, Marc W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Robert A.
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Pred. No. 1.1;
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RESULT 10
US-09-974-300-2790/c
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CURRENT FILING DATE: 2001-02-09
NUMBER OF SEQ ID NOS: 943
SOFTWARE: FastSEQ for Windows Version
LENGTH. DATE: 2001-02-09
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US-09-822-827-607
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Sequence 2790, Application US/09974300
Patent NO. US2002016721A1
GENERAL INFORMATION:
APPLICANT: Berka, Randy M.
APPLICANT: Clausen, Ib Groth
APPLICANT: Clausen, Ib Groth
TITLE OF INVENTION: Methods For Monitoring Multiple Gene
TITLE OF INVENTION: Expression
                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/822,827 CURRENT FILING DATE: 2001-03-28 NUMBER OF SEQ ID NOS: 982 SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 607 LENGTH: 22
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Best Local Similarity
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Best Local Similarity
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: McNeill, Patricia D.
APPLICANT: Houghton, Raymond L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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                                                                                                                                                                                                               1 ATGTGGGTCCCGGTTGTCTT 20
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Hepler, William
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Carter, Darrick
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Pred. No.
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Pred. No.
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ION: Expression 10085.500-US

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CURRENT APPLICATION NUMBER: US/09/974,300
CURRENT FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: 09/680,598
PRIOR FILING DATE: 2000-10-06
PRIOR PPLICATION NUMBER: 60/279,526
PRIOR FILING DATE: 2000-10-327
NUMBER OF SEO ID NOS: 8481
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2790
LENGTH: 612
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; Sequence 429, Application US/09764868
; Patent No. US20020168711A1
; GENERAL INFORMATION:
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US-09-036-613-4/c
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; ORGANISM: Bacillus licheniformis
US-09-974-300-2790
                                                                                                                                                                                                                                                                                  NAME/KEY: SITE
LOCATION: (10)
OTHER INFORMATION: n equals a,t,g, of c
NAME/KEY: SITE
LOCATION: (1410)
OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: SITE
LOCATION: (1418)
OTHER INFORMATION: n equals a,t,g, or c
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CURRENT FILING DATE: 2001-01-17
Prior application data removed - refer to PALM or file wrapper
NUMBER OF SEQ ID NOS: 1510
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 429
LENGTH: 1484
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Best Local
           Sequence 4, Application US/09036613 Patent No. US20020115135A1 GENERAL INFORMATION:
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Best Local Similarity
Matches 18; Conserv
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TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PTZ32
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OTHER INFORMATION: n equals a,t,g, or c
NAME/KEY: SITE
LOCATION: (9)
OTHER INFORMATION: n equals a,t,g, or c
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APPLICANT:
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Similarity 87.5%;
21; Conservative
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Pred. No. 3
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RESULT 13
US-09-664-761-16219
Sequence 16219, Application US/09864761
Sequence 16219, Application US/09864761
Patent NO. US20020048763A1
Patent NO. US20020048763A1
FAPLICANT: Penn, Sharron G.
APPLICANT: Hank, David K.
APPLICANT: Hankel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Acomica-X-1
CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
CURRENT FILING DATE: 2001-05-23
CURRENT FILING DATE: 2001-05-23
CURRENT FILING DATE: 2001-05-23
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
TELEPAX: 650-845-4166
TELEPAX: 650-845-4166
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 2484 base pairs
TYPE: nucleic acid
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TOPOLOGY: L...
IMBEDIATE SOURCE:
LIBRARY: BRAINOS12
LTONE: 2762136
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ETIING DATE: Filed Herewith
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
ETILING DATE:
ETILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: BILLINGS, LUCY J.
REGISTRATION NUMBER: 36,749
REGISTRATION NUMBER: PF-0
REFERRECE/DOCKET NUMBER: PF-0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSED for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Tang, Y. Tom
APPLICANT: Corley, Neil C.
TITLE OF INVENTION: SYNABS
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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CITY: Palo Alto
STATE: CA
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1 Similarity 90.0%;
18; Conservation
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linear
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APPLICATION NUMBER: PCT/US01/00664 APPLICATION NUMBER: PCT/US01/00667 APPLICATION NUMBER: PCT/US01/00666

2001-01-30 2001-01-30

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US-09-938-842A-2277/c

; Sequence 2277, Application US/09938842A

; Patent No. US20020160378A1
; TYPE: DNA
; ORGANISM: Arabidopsis thaliana
US-09-938-842A-2277
                                                                                                                             APPLICANT: Harper, Jeff
APPLICANT: Kreps, Joel
APPLICANT: Wing, Xun
APPLICANT: Wing, Xun
APPLICANT: Zhu, Tong
APPLICANT: Zhu, Tong
APPLICANT: Lor INVENTION: STRESS-REGULATED GENES OF PLA
TITLE OF INVENTION: SAME, AND METHODS OF USE
FILE REFERENCE: SCRIP1300-3
CURRENT APPLICATION NUMBER: US/09/938,842A
CURRENT FILING DATE: 2001-08-24
PRIOR APPLICATION NUMBER: US 60/227,866
PRIOR FILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: US 60/264,647
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR APPLICATION UNMBER: US 60/300,111
PRIOR APPLICATION NUMBER: US 60/300,111
PRIOR FILING DATE: 2001-06-22
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: OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL US-09-864-761-16219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 16219
LENGTH: 569
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                                                                                                            PRIOR FILING DATE: 200 NUMBER OF SEQ ID NOS:
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                                                                   ENGTH: 975
                                                                                           ID NO 227
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 60/234,687 FILING DATE: 2000-09-21 APPLICATION NUMBER: US 09/608,408 FILING DATE: 2000-06-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/US01/00662
FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00661
FILING DATE: 2001-01-30
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FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00663
FILING DATE: 2001-01-30
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85.7%;
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Pred. No. 89;
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Search completed: November 19, 2002, 10:16:24 Job time : 90.2308 secs
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US-09-814-950-3
                                                                                                                                                                                                                                                                                            SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/09814950 Patent No. US20020137130A1
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Best Local Similarity
                                                                                                                  Matches
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CURRENT FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: I TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: CL001174
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: WEI, Ming-Hui et al
                                                                                                                                                                                                                                                                                                         SOFTWARE:
                                                         27025 TGTGGGTCCCAGGTGCCTTCC
                                                                                                                                                                                                     NAME/KEY: misc_feature LOCATION: (1)...(33239)
                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                     OTHER INFORMATION: n =
                                                                                                                                                                                                                                    FEATURE
                                                                                                                                                                                                                                                                             ENGTH:
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85.7%;
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Pred. No. 9
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                                                                                                                                            DB 10; | Length 33239;
                                                                                                                                                                                                                                                                                                                                                                                            PROTEINS, ENCODING HUMAN SECRETED PROTEINS, AND
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GenCore version 5.1.3 Copyright (c) 1993 - 2002 Compugen Ltd.

OM nucleic - nucleic search, using sw model November 19, 2002, 00:33:44; Search time 1026.35 Seconds (without alignments) 765.605 Million cell updates/sec

Title: Perfect score:

Run on:

Sequence: US-09-829-004A-8 27 1 gttctggtgcacccccagtgggtcctc 27

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 2054640 seqs, 14551402878 residues

Total number of hits satisfying chosen parameters: 4109280

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

GenEmbl:*

gb_ba:*
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

RESULT 1 AXXB1251 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL	C 22 22 22 22 22 23 33 33 33 33 33 33 33	Result No. S
	888 - 888777777777777777777777777777777	core
AX281251 Sequence 8 AX281251 AX281251.1 Synthetic synthetic artificial pedyczak, A Immunogeni and uses t Patent: WO	######################################	ry
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linear PAT 02-NOV-200	AX281251 Sequence AX281251 Sequence XI3941 Human PSA 9 X75755 PSA=prostat M21897 Human prost AJ310937 Homo sapi AR153342 Sequence BD007601 Method fo AJ459782 Homo sapi U17040 Human prost I67863 Sequence AX267642 Sequence AX267642 Sequence AX267642 Sequence AX267642 Sequence AX267643 Homo sapi AF335477 Homo sapi AF335477 Homo sapi AF335478 Homo sapi AZ300986 Sequence AX267642 Sequence AX267642 Sequence AX267643 Homo sapi AJ31896 Human mRNA BC005307 Homo sapi X73560 M. mulatta m AJ310938 Homo sapi X73560 M. mulatta m AJ310938 Homo sapi X73560 Sequence AX0337560 Sequence X07730 Human mRNA BC005307 Homo sapi AX106218 Sequence X07330 Human mRNA BC005307 Homo sapi AX106218 Sequence X0733404 Sequence X0733407 Sequence X0733407 Sequence AX033407 Sequence AX033408 Sequence AX033408 Sequence AX033407 Human prost AC027602 Homo sapi AC027603 Homo sapi AC027603 Homo sapi AC027603 Homo sapi AC027604 Homo sapi AC027604 Homo sapi AC027605 Sequence AR105688 Sequence AR105695 Sequence AR105695 Sequence AR105695 Sequence AR105695 Sequence AR105695 Sequence	; #

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REFERENCE
AUTHORS
TITLE
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HSPSA2
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FEATURES
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AR047835
LOCUS
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SOURCE
ORGANISM
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ACCESSION
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Matches 27
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                                                                                                                                                                                                                                                                                                                      Direct Submission
Submitted (09-JAN-1989) Digby M.R., The Howard Florey Institute,
Melbourne University, Parkville; Victoria 3052, Australia
(bases 1 to 287)
Digby, M., Zhang, X.Y. and Richards, R.I.
Human prostate specific antigen (PSA) gene: structure and linkag,
to the kallikrein-like gene, hGK-1
Nucleic Acids Res. 17 (5), 2137 (1989)
89183632
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1 (bases 1 to 200)
Gundling,GJ.
Rapid RNA isolation procedure in the presence
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Sequence 1 from patent US
AR047835
AR047835.1 GI:5969300
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Similarity 100.0%;
27; Conservative 0.
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X13941
X13941.1 GI:35724
kallikrein; prosta
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Location/Qualifiers
1. .200
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                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
1. (bases 1 to 287)
                                                                                                                                                                                         Homo sapiens
                                                                                                                                          Digby, M.R.
                                                                                                                                                                                                                                                                                                                                                                    Conservative
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kindly reviewed (16-may-1989) by Digby M.R. Location/Qualifiers
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PSA gene for prostate specific antigen
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Location/Qualifiers
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/db_xref="taxon:32630"
/note="CLP314"
10 c 8 g 7 t
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58 c 67 g
                                                                                                                                                                                                               prostate specific antigen; PSA gene
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Pred. No.
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                                                                                                                                                                              Euteleostomi;
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95079406
7527295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens breast cancer specimen.
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PSA-prostate-specific antigen
Partial, 569 nt].
$75755
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 569)
Monne, M., Croce, C.M., Yu, H. and Diamandis, E.P.
Molecular characterization of prostate-specific antigen messenger
RNA expressed in breast tumors
Cancer Res. 54 (24), 6344-6347 (1994)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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27; Conservative
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/Ob_xref="taxon:9606"
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/Clone_lib="cosmid"
<1. 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /protein_id="CAA32124.1"
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/db_xref="SWISS-PROT:PO7288"
/translation="ALRESSCLIGHWEAGSARSIPNEGRCLWPLVAGQSAAVFWCTPSG
SSQLPTASG"
268. >287
/note="intron II"
100 c 78 g 56 t
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/codon_start=3
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EKHSQPWQVLVASRGRAVCGGVLVHPQWVLTAAHCIRNKSVILLGRHSLFHPEDTGQV
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                                                                                                                            /gene="PSA"
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                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                               'gene="PSA"
                                                                                                                                                                                                                                                    'partial
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[human, breast cancer
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specimen, mRNA
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RESULT 6
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 HSA310937
Homo sapiens mRNA
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88326297
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Riegman, P. H., Klaassen, P., van der Korput, J.A., Romijn, J.C. and
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Glycoprotein; prostate antigen; prostate-specific antigen;
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                                                                                                                                                                                               /note="prostate specific antigen" 5 a 195 c 206 g 132 t upstream of PvuII site.
                                                                                                                                                                                                                                                     /gene="APS"
/note="prostate specific antigen signal peptide"
62 . .559
                                                                                                                                                                                                                                                                                                             /db_xref="GDB:G00-119-695"
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VCGGVLVHPOWVLTAAHCIRNKSVILLGRHSLFHPEDTGQVFQVSHSFPHPLYDMSLL
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CTPGPDGAAGSPDAWV"
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182 c 160 g 119 t
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                                                                                                                                                                                                                                                                                                                                                                                                                     /gene="APS"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /gene="APS"
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708 bp for putative p
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preproPSA-RP2 (KLK3 gene),
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(PA) gene,
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3' end,
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clone PA 424.
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Matches 27; Conserv
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Sequence 4
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1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
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AJ310937.1 GI:14422304
alternative splicing; HKLK3 gene; kallikrein 3; preproPSA-RP2.
AR153342.1 GI:15120874
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Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                               /gene="KLK3"
/evidence=experimental
/ 214 c 219 g
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<1. .37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /product="PSA-RP2"
578. .708
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110. .577
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MSLLKNRFLRSPGDDSSHDLMLLRLSEPAELTDAVKVMDLFTQEFALGTTCYASGWGSI
MSLLKNRFLRSPGDDSSHDLMLLRLSEPAELTDAVKVMDLFTQEFALGTTCYASGWGSI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /gene="KLK3"
/function="unknown"
/note="alternative splicing, transcript
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/clone="NH101007.97"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'gene="KLK3"
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                                         711 bp
from patent US 6235486.
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ORGANISM
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                                                                                                                                                                        Query Match
Best Local S
Matches 27
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JOURNAL
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Best Local
Matches 2
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E 1 (bases 1 to 711)

S Tindall,D.J., Young,C.Y.F., Mccormic,D.J., Klee,G.G., Sae Kumar,A., Rittenhouse,H.G. and Wolfert,R.L.

Method for detecting metastatic prostatic cancer patent: JP 2001503991-A 14 27-MAR-2001;

MAYO FOUNDATION FOR MEDICAL EDUCATION AND RESEARCH,HYBRIT OS Unidentified PN JP 2001503991-A/14

PD 27-MAR-2001

PD 27-MAR-2001

PR 14-NOV-1997 JP 1998522888

PR 14-NOV-1996 US 08/759354

PR 14-NOV-1997 JF 11998522888

PR 14-NOV-1997 JF 110DALL,CHARLES Y F YOUNG,DANIEL J MCCORMIC
                                                                                                            GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                       GTTCTGGTGCACCCCAGTGGGTCCTC
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1 (bases 1 to 711)
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2 (constant)
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PC C1
G01N33,
PC G6
CC S1
CC T0
FH K6
FT S0
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PI MC
                                                                                                                                                                             Similarity
27; Conser
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JP 2001503991-A/14.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     unidentified.
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JP 2001503991-A/14
27-MAR-2001
14-NOV-1997 JP 1998522888
14-NOV-1996 US 08/759354
DONALD J TINDALL, CHARLES Y F YOUNG, DANIEL J MCCORMIC, PI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C12N15/09, C07K16/40, C12N5/10, C12N9/64, C12P21/08, C12Q1/68,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GE G KLEE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 27; Dilarity 100.0%; Pred. No. 0. Conservative 0; Mismatches
                                                                                                                                                                        100.0%; llarity 100.0%; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Topology: Linear;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Strandedness: Double;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          G01N33/574,C12N15/00,C12N5/00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OHAMMAD SAEED SAEDI,ABHAY KUMAR,HARRY G RITTENHOUSE, PI
L WOLFERT
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for detecting metastatic
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                                                                                                                                                                                                                                                                                                                                                   /organism='Unidentified'.
Location/Qualifiers
1. .711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="unknown"
208 c 206 g
                                                                                                                                                                                                                                                                                    /organism="unidentified"
/db_xref="taxon:32644"
208 c 206 g 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                 Score 27;
Pred. No.
                                                                                                                                                                             Mismatches
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0.049;
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0.049;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-JAN-2002
                                                                                                                                                                             Gaps
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VERSION
KEYWORDS
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HSA459782
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       REFERENCE
                                                                                                                                                                                                    DEFINITION
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TITLE
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Best Local (
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                                                                   ORGANISM
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Courty,Y.

Direct Submission
Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire d'Enzymologie, Universite F. Rabelais, 2 bis byd Tonnelle, 37032 cedex, FRANCE
37032 cedex, FRANCE
                                                                                                                                                                                                                                                                                                                                                l Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             alternative splice variant sequences: Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Heuze-Vourc'h,N. and Courty,Y. Complex alternative splicing of the hKLK3 gene coding for the tumour marker PSA (prostate-specific-antigen) unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
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Homo sapiens
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                    Homo sapiens variant 1.
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                                                                          alternative splicing; KLK3 gene;
                                                                                            АJ459782.1
                                                                                                                                                                         HSA459782
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                             185
                                                                                                                                                                                                                                                                                                                                                100.0%;
llarity 100.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /gene="KI
94..804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /translation="mwvpvvfltlsvtwigaaplilsrivggwecekhsqpmqvlvas rgravcggvlvhrgwrtjaahcltnksvtllghsledjagvergvshsphplyd msllknrelrpgddsshdlmllrlspapaeltdavkvwblptgopalgttcasgwesi epeepltpkklqcvblhvlsndvcagvhpqkvtkfmlcagrafggsstcsdsggbly cggvlqgitswgsepcalperpslytkvvhyrkwikdtivanp"
                                                                                                                                                                                                                                                                                                                                                                                                                                        /product="prostate specific antigen" 257 c 242 g 186 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /product="prostate specific antigen"
/protein_id="CAD30845.1"
/db_xref="GI:20520643"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="YC140405-00"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'gene="KLK3"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 evidence=experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'gene="KLK3"
'note="alternative splice variant
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22. .807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /tissue_type="prostate"
L. .870
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                                                                                                                                                     mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "KLK3"
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                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                   876 bp
                                                                                                                                                                                                                                                                                                                                                Score 27; DB Pred. No. 0.0 0; Mismatches
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specific
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                                                                          prostate specific antigen
                                                                                                                                                                                                                                                                                                                                                                  DB 9;
0.049;
                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                     linear
antigen (
                                                                                                                                                                                                                                                                                                                                                                                    Length 870;
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                                                                                                                                                                                                                                                                                                                                                   Indels
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(KLK3 gene), splice
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JOURNAL
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SOURCE
ORGANISM
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HSU17040
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Best Local S
Matches 27
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Human prostate specific antigen precursor mRNA, complete cds.
U17040
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Submitted (03-MAY-2002) Courty Y., EMI-U 0010, Laboratoire
d'Enzymologie, Universite F. Rabelais, 2 bis bvd Tonnelle,
                                                 Submitted (08-NOV-1994) Jose M J G M Jefferson University Hospital, 1025
                                                                                                                                              2 (bases 1 to 990)
Moreno, J.M.
                                                                                                                                                                                                      A Comparative Analysis of Prostate Specific Antigen Gene Sequence
in Benign and Malignant Prostate Tissue
Unpublished
                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 990)
                                                                                                               Direct Submission
                                                                                                                                                                                                                                                                                                Gomella, L.G.
                                                                                                                                                                                                                                                                                                                            Monne, M.M., Moreno, J.M., Mele, C.M.; Mulholland, G.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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Complex alternative splicing of the hKLK3 gene coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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/gene="KLK3".
94. .681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /product="prostate specific antigen"
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/translation="MAVPVVFLTLSVT"
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/translation="MAVPVVFLT"
/translation="MAVPVVFLT"
/translation="MAVPVVFLT"
/translation="MAVPVVFLT"
/translation="MAVPVVFLT"
/translation="MAVPVVFLT
Location/Qualifiers
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22. .684
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/db_xref="taxon:9606"
/clone="YC171105-00"
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235 c 231 g 190 t
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                                                    J G Moreno, U
1025 Walnut 9
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0.049;
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Street, P
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                                                       Philadelphia, PA
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                       1130 bp
Homo sapiens prostate-specific a
cds, alternatively spliced.
AF335477
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Croce, C., Gomella, L., Mulholland, S.Grant.,
Fischer, R.
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Sequence 13 from patent US 5674682.
167863
167863.1 GI:2829985
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              AF335477.1
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                                                                                                                                                                                                                                                                                     Patent:
                                                                                                                                                                                                                                                                                                  cancer
                                                                                                                                                                                                                                                                                                            Nucleic acid primers for detecting micrometastasis of prostate
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93. .113
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114. .8
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EPEEFLTPKKLQCVDLHVISNDVCAQVHPQKVTKFMLCAGRWTGGKSTCSGDSGGPLV
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                                                                                                                                                                                                                                  /organism="unknown"
291 c 272 g
                                                                                                                                                                                                                                                                      Location/Qualifiers
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291 c 273 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /translation="MWVPVVFLTLSVTWIGAAPLILSRIVGGWECEKHSQPWQVLVAS
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                                                                                                                                                                        0
                                                                                                                                                                                  Score 27;
Pred. No.
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                                               linear PRI 13-MAY-
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                                                            PRI 13-MAY-2002
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MEDLINE
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Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Butheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1130)

David,A., Mabjeesh,N., Azar,I., Biton,S., Engel,S., Bernstein,J.,

Romano,J., Avidor,Y., Waks,T., Eshhar,Z., Langer,S.Z.,

Lifschitz-Mercer,B., Matzkin,H., Rotman,G., Toporik,A., Savitsky,K.

and Mintz,L.

Unusual alternative splicing within the human kallikrein genes KLK2

and KLK3 gives rise to novel prostate-specific proteins
J. Biol. Chem. 277 (20), 18084-18090 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ch 100.0%; Score 27; DB 9; Length 1130; 1 Similarity 100.0%; Pred. No. 0.09; Length 1130; 27; Conservative 0; Mismatches 0; Indels
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Submitted (11-JAN-2001) Compugen Ltd.,
69512, Israel
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11834722
                                                                                                       Patent: WO 0151633-A 616 19-JUL-2001; CORIXA CORPORATION (US)
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vid, A., Engel, S., Azar, I., Bernstein, J.,
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                284
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PSpDRELGSPLSLPAPLQAHTPSPSILQQSSLPHQVPAPSHLPQNFLPIAQPAPCSQL
                                                                    Location/Qualifiers
1. .1350
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/chromosome="19"
/map="19q13.3"
              /organism="Homo sapiens"
/db_xref="taxon:9606"
404 c 391 g 27
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                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                                              Search completed: November 19, Job time: 1028.68 secs
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                                                                                                                        GTTCTGGTGCACCCCCAGTGGGTCCTC 804
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Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                            Patent: WO 0173032-A 616 04-OCT-2001; CORIXA CORPORATION (US)
                                                                                                                                                                                                                                                                                                                                                                   Xu,J., Dillon,D.C., Mitcham,J.L., Harlocker,S.L., Jiang,Y., Kalos,M.D., Fanger,G.R., Retter,M.W., Stolk,J.A., Day,C.H., Vedvick,T.S., Carter,D., Li,S.X., Wang,A., Skelky,Y.A., Hepler,W.T. and Henderson,R.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AX267642.1 GI:16516315
                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
404 c 391 g 27
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1. .1350
                                                                 2002, 04:02:54
3.15
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                                                                                                                                                                                 Gaps
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5.1.3

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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                                                                                                                                                                 Result
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Database :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                   ŏ
                                7654627
                                                                                                                                                                                                                        Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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BC
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seq
   N_Geneseq_101002:*
1: /SIDS2/gcgdata/gc2: /SIDS2/gcgdata/gc3: /SIDS2/gcgdata/gc4: /SIDS2/gcgdata/gc5: /SIDS2/gcgdata/gc5: /SIDS2/gcgdata/gc6
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Match
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Gapop 10.0 , Gapext 1.0
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27
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                                                                                                                                                                                                                                                                                      /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1989.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1990.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1991.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1992.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1993.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1994.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1994.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1994.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1995.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1997.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1997.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1999.DAT:
/SIDS2/gcgdata/geneseq/geneseqn-emb1/NA1999.DAT:
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                 00000000
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1982.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1983.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1984.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1984.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1985.DAT:*
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2001A.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2001B.DAT:*
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2002.DAT:*
                                                                                                                                                   Length
              178
178
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208
517
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               AAT92975
AAA61482
AAH41845
AAX08947
                                                                        AAD21362
ABA92483
ABA01120
AAV60379
                                                                                                                                                                                             SUMMARIES
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          DNA oligonucleotid
Human PSA nucleoti
mRNA target sequen
cDNA for prostate
prostate specific
Prostate specific
Human prostate spe
                                                                                                                                           Description
cDNA encoding huma
                                                                                                                     Prostate specific
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AAD21362
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27 100 0 135 27 100 0 135 27 100 0 144 27 100 0 146 27 100 0 146 27 100 0 146 27 100 0 172 27 100 0 172 27 100 0 172 27 100 0 172 27 100 0 210 27 100 0 210 27 100 0 342 27 100 0 342 27 100 0 466 27 100 0 466 27 100 0 466 27 100 0 172 27 100 0 172 27 100 1 186 27 100 1 186 28 94 1 186 29 1 1 186 21 1 186 22 1 1 186 23 8 88 1 23 23 8 88 1 23 23 8 88 1 27 23 8 88 1 27	7 100.0 7 100.0 7 100.0 7 100.0 7 100.0 7 100.0 7 100.0
1 1 2 2 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3	0 23 0 23 0 23 2 20 2 22 9 23 9 23
AAS63953 AAS63953 AAS93860 ABL95324 AAV22497 AAV22496 AAA11711 AAV3249667 AA71048667 AA7104867 AAF69732 A	ABV25078 AAZ47134 AAT91054 AAX99846 AAH99720 AAC03740 ABV23473 ABV29201
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Human proprostate Prostate Human proprostate Human proprostate Prostate Human wo DNA enco DNA enco DNA enco DNA enco DNA enco Human pr Single n Human se Human se Human se	Human Human Human Prost Human Human Human

ALIGNMENTS

AAD21362 standard; DNA;

27 ВР

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Prostate-specific antigen; PSA; immunogenic peptide; immune response; therapy; cancer; prostate cancer; tumour metastasis; prophylaxis; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAD21362;
(AVET ) AVENTIS PASTEUR LTD
                                            10-APR-2000; 2000US-195456P
                                                                                         10-APR-2001; 2001WO-CA00473
                                                                                                                                      18-OCT-2001.
                                                                                                                                                                                  WO200176622-A2
                                                                                                                                                                                                                                                                                                                                                                                      Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prostate specific antigen (PSA) derived peptide, CLP314 encoding DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                      /product= "PSA derived peptide,
/note= "CDS does not include st
/partial
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1..27
                                                                                                                                                                                                                                                    start and
                                                                                                                                                                                                                                                                       CLP314"
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2001-663015/76

A,

Chong P,

CDY;

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RESULT 2
ABA92483
ID ABA92483
ID ABA92483
AXX ABA9
AXX ABA9
AXX DETE
AXX WPI
AXX TELL
AXX T
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Best Local S
Matches 27
The present invention describes a method for detecting the occurrence of a nucleic acid synthesising reaction during the synthesis of a nucleic acid with an enzyme. The method comprises the use of an insoluble substance formed in the reaction as an indication. The method can be used for detecting the presence or absence of a nucleic acid synthesising reaction (especially a nucleic acid amplification reaction). The method is useful in gene analysis, diagnosis of cancer and other diseases, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The patent discloses immunogenic peptides of prostate-specific antigen (PSA) and nucleic acids encoding them. The peptides of the invention are used to prepare a medicament to elicit an immune response in an animal. They are used to treat cancer such as prostate cancer and tumour metastasis. They are also useful for prophylaxis, for preparing monoclonal or polyclonal antibodies, and in conventional techniques of immunology, molecular biology, cell biology and recombinant DNA technology. The present sequence is a DNA encoding PSA derived peptide, CLP314.
                                                                                                                                                                                                                                                                                                  Detecting the product of an enzyme-based nucleic acid synthesizing reaction, useful in gene analysis, diagnosis of cancer and other diseases, and in identifying bacteria, comprises the formation of an insoluble substance -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mori
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection;
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                                                                                                                                                                                                                                         Example 4; Page 18; 44pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-075170/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-MAY-2000; 2000JP-0132667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-APR-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (EIKE )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EIKEN KAGAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nagamine K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleic acid synthesis; identification; cancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 10 C; 8 G; 7 T; 0 other;
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Pred. No. 0.0
0; Mismatches
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0.029;
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RESULT 4
AAV60379
ID AAV

AAV60379 standard; mRNA;

200

ВP

Qy

ш

GTTCTGGTGCACCCCCAGTGGGTCCTC

GTTCTGGTGCACCCCCAGTGGGTCCTC

Query Match

Local

Similarity

100.0%;

Score 27; Pred. No. Mismatches 27 65

DB 24; 0.034;

Length 178;

0;

0;

Gaps

0;

Conservative

0,

Sequence 178 BP;

33 A; 56 C; 50 G; 39 T; 0 other;

DЪ

39

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DЬ
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                  The invention relates to a method for synthesising a nucleic acid using a double-stranded nucleic acid as template and incubating under conditions allowing the establishment of a complementary chain synthesis reaction. The method uses an arbitrary primer to initiate the complementary chain synthesis reaction. The method is particularly useful in gene and disease diagnosis. It is a highly efficient and reaction specific method in which no temperature variation is required. The present sequence is used in an example illustrating the invention.
                                                                                                                                                                 Isothermal amplification of nucleic acids using double-stranded nucleic acid as template to establish complementary chain synthesis reaction from primer enabling base pairing in domain to be annealed, useful e.g. in gene diagnosis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 178 BP; 33 A; 56 C; 50 G; 39 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           in identifying bacteria. The present sequence oligonucleotide used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA01120;
                                                                                                                                    Example 1; Page 63; 75pp; Japanese.
                                                                                                                                                                                                                            WPI; 2002-010907/01
                                                                                                                                                                                                                                                   Notomi T,
                                                                                                                                                                                                                                                                                                   07-APR-2000; 2000JP-0111939
                                                                                                                                                                                                                                                                                                                            30-MAR-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                PSA nucleotide sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid synthesis; complementary chain synthesis; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                   Nagamine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                            2001WO-JP02771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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RESULT 5
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                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                     Best
                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a mRNA target sequence from exons 2 and 3 of the prostate specific antigen (PSA) gene... The sequence is used to exemplify the invention. The specification describes a method for purifying RNA. The method comprises contacting a test sample with polyvalent transition metal ions to form a precipitant and a supernatant, separating the precipitant from the supernatant, and collecting the supernatant to obtain a purified solution of total RNA. The RNA is used for nucleic acid amplification and disease detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Purification of RNA, used for nucleic acid amplification and disease detection - by precipitating other sample components with transition metal ions, separating and collecting the purified RNA
                                                                                      Diagnosis; prostate cancer; prostate specific antigen; PSA; metastatic prostate cancer; prostate cell; secondary tumour; bone metastatic anchoring; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Columns 9-10; 6pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA
16-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gundling
                      23-OCT-1997.
                                           WO9739139-A1
                                                                                                                                   cDNA for prostate
                                                                                                                                                                                                      AAT92975 standard; cDNA; 208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-556473/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-OCT-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-DEC-1998
                                                                   Homo
                                                                                                                                                           24-APR-1998
                                                                                                                                                                                  AAT92975
                                                                                                                                                                                                                                                                                                                                                     Sequence 200 BP; 31 A; 58 C; 67 G; 44 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ABBO ) ABBOTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV60379;
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                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                        1 GTTCTGGTGCACCCCCAGTGGGTCCTC
                                                                 sapiens
                                                                                                                                                                                                                                                            GTTCTGGTGCACCCCCAGTGGGTCCTC 146
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              target sequence;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    target
                                                                                                                                                                                                                                                                                                          l Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                          (first entry)
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97WO-US06497.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LAB.
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                                                                                                                                    specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    from exons 2
                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prostate specific antigen; PSA; RNA purification;
                                                                                                                                    antigen
                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                     Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                   27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and 3 of PSA gene
                                                                                                                                    (PSA).
                                                                                                                                                                                                                                                                                                                     0.035;
                                                                                                                                                                                                                                                                                                                              DB 19;
                                                                                                                                                                                                                                                                                                                                                     0 other;
                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                200;
                                                                                                                                                                                                                                                                                                          0;
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RESULT 6
AAA61482
Db
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cDNA sequences AAT92974-77 encode the prostate specific antigen (PSA). The detection of PSA mRNA in peripheral blood is associated with metastatic prostate cancer. Detection of PSA mRNA in the bloodstream indicates that prostate cells are circulating in the blood and confirms diagnosis of cancer. Circulating prostate cells also indicate a risk of a secondary tumour or bone metastatic anchoring. Detection of the PSA mRNA after prostatectomy or radiotherapy indicates a risk that the prostate cancer has spread and that the surgery and radiotherapy was not effective. PCR primers AAT92971-73 were used for the monitoring and diagnosis of patients with prostate cancer. The method comprises the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Monitoring or diagnosis of prostate cancer - by detecting prostate-specific antigen mRNA using specific primers
                                                                                                                                                                                                                                                                                                                                                        AAA61482 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnosis of patients with prostate cancer. The method comprises the detection of PSA mRNA by reverse transcriptase polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure;
                                                                                                                                                                                                                                                                                        Prostate
                                                                                                                                                                                                                                                                                                             23-OCT-2000
                                                                                                                                                                                                                                                                                                                                   AAA61482;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                                     GTTCTGGTGCACCCCCAGTGGGTCCTC
                                                                                                                                                                                                                                                                                                                                                                                                               GTTCTGGTGCACCCCCAGTGGGTCCTC 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
27; Conser
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                                                                                                                                                                                                                                                                                       specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29
                                                                                                                                                                                                                                                                                       antigen gene
                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 62 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25pp;
                                                                                                                                                                                                                                                                                                                                                         517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        English.
                                                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     73 G; 44 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 27;
                                                                                                                                                                                                                                                                                       fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                     27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .035;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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Homo sapiens
                                                      Isothermal DNA amplification; nucleic acid synthesis; PSA gene template; prostate specific antigen; human; c
                                                            ds
                                                                                          detection;
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Example

8

Fig 17;

95pp; Japanese

Nucleic acid synthesis using primer containing a one on the template for isothermal amplification

sequence identical using simple reager

reagents

WPI; 2000-376587/32

Notomi

Ţ,

Hase T;

09-NOV-1998; 08-NOV-1999;

98JP-0317476 99WO-JP06213

(EIKE) EIKEN KAGAKU KK.

18-MAY-2000

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RESULT ANAHALISA NE SULT NA CANANA NE SULT NA CA
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A region is produced at the 3 end of a target sequence which contains sub-sequences designated F1, F2C and F1C, where F1C is the complement of F1. A loop is formed by annealing of F1 to its complementary sequence F1C, and a primer containing a sub-sequence F2 (complementary sequence F1C, and a primer containing a sub-sequence F2 (complementary strand to the template is synthesised using a DNA polymerase, and the process is C2 repeated using sub-sequences at the 3' end of the complementary strand designated R1, R2C and R1C, using a primer containing R2 at its 3' end. C3 designated R1, R2C and R1C, using a primer containing R2 at its 3' end. C4 cand methods for the detection of a nucleic acid sequence in a sample C4 cand methods for the detection of a nucleic acid sequence in a sample C5 cand methods for the detection of a sequence complementary to the loop C6 and methods for the detection of a sequence complementary to the loop C6 can efficient and highly specific method of the invention provides C6 of a target nucleic acid sequence using a simple reagent mixture. C6 The present sequence represents a prostate specific antigen (P5A) C7 gene fragment used as a template for the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 27
The resent invention describes a new method for synthesising nucleic acids comprising: (a) preparing a nucleic acid, with an F2c region, able to form a loop by annealing the F1 region of the 3' end to the F1c region on the same chain; (b) synthesis of a complementary chain using the 3' end of F1 as the starting point; (c) replacing the complementary to F2c) at its 3' end and using it as a starting point for complementary chain synthesis by a polymerase; and (d) replacing the chain synthesised in step c by annealing a polynucleotide containing a region complementary to a part of the chain and using its 3' end as the starting point. Also described are: (1) a kit for synthesising nucleic acids; and (2) a kit for detecting target base sequences. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 8; Fig 17; 96pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthesizing nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Notomi T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EIKE ) EIKEN KAGAKU KK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-NOV-1999; 99WO-JP06213
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-MAY-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prostate specific antigen; nucleic acid synthesis; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prostate specific antigen related nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH41845;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001-343603/36.
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27; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           517 BP; 94 A; 163 C; 151 G; 109 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hase T;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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0.038;
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RESULT 8
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Best Local s
Matches 27
                                  Human kallikrein 2 (hK2) is expressed at elevated levels relative to the prostate cancer antigen, prostate specific antigen (PSA) by breast cancer cells when stimulated by androgens. Detecting levels of hK2 may provide a simple diagnostic tool for detecting or determining breast cancer. Detecting hK2 is achieved by producing an hK2 DNA from hK2 RNA by reverse transcription. The hK2 DNA obtained is then amplified by PCR and detected using probes. Determination of breast cancer may also be determined by exposing the hK2 polypeptide to an agent which binds to it and then detecting the binary complex formed. The amount of complex formed correlates directly to the presence of breast cancer in the human individual. The methods can be used in the diagnosis, treatment and/or monitoring of the progression or remission of breast cancer and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sequence represents a prostate specific antigen related sequence, which is used in an example from the present is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 517 BP; 94
                                                                                                                                                                                                                                          Disclosure; Page 34-35;
                                                                                                                                                                                                                                                                      Use of human kallikrein 2 - as a marker for developing products for the diagnosis, prognosis, monitoring and treatment of breast cancer
                                                                                                                                                                                                                                                                                                                                                             Klee GG,
                                                                                                                                                                                                                                                                                                                                                                                       (KLEE/) KLEE G G.
(MAYO-) MAYO FOUNDATION.
(TIND/) TINDALL D J.
(YOUN/) YOUNG C Y F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human kallikrein 2; hK2; breast cancer; diagnosis; detectitreatment; monitoring; prostate specific antigen; PSA; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9859073-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human prostate specific antigen (PSA) gene.
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l similarity 100.0%;
27; Conservative 0
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1..711
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Prostate_specific_antigen
                                                                                                                                                                                                                                           67pp; English.
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                                                                                                                                                                                                                                                                                                                                                               CYF;
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Pred. No. 0.038;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        517;
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monitoring of the progression monitoring hK2 levels.

Sequence 711 BP; 148 A;

208 c;

206

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149 T;

0 other

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RESULT 9
ABK86205
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The invention relates to a method of inducing a cell-mediated immune cresponse against a cell comprising a target antigen (I) in a subject, treating a subject having undesired cells, for example tumour cells or virally infected cells (C), reducing the number of (C) in a subject, and stimulating presentation of (I) by a cell. This is done by administering a polynucleotide (II) exoding a variant of (I), so that (II) expressed in a cell and cell-mediated immune response is induced. The method can be used to treat prostate cancer, breast cancer, colorectal cancer and pancreatic cancer, as well as lymphomas and leukaemias. The method is also useful in treating chronic viral infections such as those caused by hepadnaviruses, lentiviruses (including human immunodeficiency virus (HIV)), herpesviruses and the flaviviruses and pestiviruses. The present sequence represents the coding sequence of human prostate specific antigen (PSA) variant which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local
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                                                                                                                                                                                                                                                  a cell,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; prostate specific antigen; PSA; cytostatic; antiviral; immunostimulant; cell-mediated immune response; tumour; breast crivirus infection; prostate cancer; colorectal cancer; pancreatic (lymphoma; leukaemia; hepadnavirus; lentivirus; herpesvirus; human immunodeficiency virus; HIV; flavivirus; pestivirus; gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA
                                                                                                                                                                                                          Disclosure; Page 121-122; 146pp; English.
                                                                                                                                                                                                                                                                             Inducing
                                                                                                                                                                                                                                                                                                                                             Mincheff MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200240059-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                                                                                                                                                                                                                                                             reducing
                                                                                                                                                                                                                                                                                                                                                                                                  (MINC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                         (ZOUB/)
                                                                                                                                                                                                                                                                                                                                                                                       (AMBI-)
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                                                                                                                                                                                                                                                                                                                                                                      ) MINCHEFF M S.
) LOUKINOV D I.
) ZOUBAK S.
                                                                                                                                                                                                                                                                                                                                                                                                             AMERICAN FOUND BIOLOGICAL
                                                                                                                                                                                                                                               g a cell-mediated immune response against a target g undesired cells and stimulating presentation of a comprises administering a polynucleotide encoding
                                                                                                                                                                                                                                                                                                         AAU98921.
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                                                                                                                                                                                                                                                                                                                                             Loukinov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
7..744
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
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                                                                                                                                                                                                                                                                                                                                             DI,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Prostate specific antigen (PSA) variant with signal sequence deleted"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                             Zoubak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    specific antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PSA) variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
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                                                                                                                                                                                                                                                  an antigen
y a variant
                                                                                                                                                                                                                                                                             antigen
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RESULT 10
ABV25078
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                       a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) or the specification or its complement. (I) is useful for:
(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                    Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, use for detecting presence of prostate cancer, stage of prostate cancer
                    patient;
(I) is a
                                                                  cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) selecting a composition for inhibiting prostate cancer in a patient;
(f) assessing the prostate cell carcinogenic potential of a compound;
                                                                                                                                                                            The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the
                                                                                                                                                                                                                Claim 1;
                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                                                                                                                           Schlegel
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18-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                          16-MAR-2000;
25-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                  17-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                (MILL-) MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       the signal sequence deleted, used the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      prostate expression marker cDNA 25069
                       also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                          ₽,
                                                                                                                                                                                                                Page 4874;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  744 BP;
                                                                                                                                                                                                                                                                                                                                                        2000US-211314P.
2000US-219007P.
2000US-255281P.
                       useful
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2000US-207454P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     marker;
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                    as a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic;
gene; ss.
                      pharmacodyanamic or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            841
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 222 C;
                                                                                                                                                                                                                                                                                                           Monahan
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                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  213 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                           JE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                carcinogen;
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                      pharmacogenomic marker
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useful
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Query Match

Sequence

841

BP;

144

A;

262 C;

262 G; 173

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Score 27;

DB

23;

Length

841;

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RESULT 11
ARAZ47134
ID ARAZ47
AC ARZ47
AC ARZ47
AC ARZ47
AX ARZ47
AX Huma
XX Huma
XX Hom
XX Gytt
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                                                                                                                               RESULT 12
AAT91054
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                                                                                                                                                                                        В
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   Best Loc
Matches
                                                                                                                                                                                                                                                             Best Local
                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                              The invention relates to a method of immunization against prostatic cancer (PC) by administering a nucleic acid (this sequence) that encodes prostate-specific antigen (PSA), or its immunologically active fragment, operably linked to regulatory sequences. The nucleic acid is taken up by cells and then expressed to generate an immune response against PSA. The method is used to protect against, or treat, PC, or more generally to induce a cellular immune response to PSA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ47134 standard; DNA; 990 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunization; prostate cancer; prostate-specific antigen; PSA;
cytostatic; immune response; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human prostate-specific antigen gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-MAR-2000 (first entry)
Prostate specific antigen; primer; human; prostate cancer;
                          Human prostate specific antigen gene.
                                                                                     AAT91054;
                                                                                                                AAT91054 standard; DNA; 992 BP
                                                                                                                                                                                                                                                                                                       Sequence 990 BP; 209 A; 291 C; 273 G; 217 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 42; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunization against prostatic cancer using nucleic acid encoding prostate-specific antigen, for treatment or prophylaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-072551/06.
P-PSDB; AAY56048.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Weiner DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-MAY-1999;
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                                                       27-FEB-1998 (first entry)
                                                                                                                                                                                      218 GTTCTGGTGCACCCCCAGTGGGTCCTC 244
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                                                                                                                                                                                                                                               l Similarity
27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kim JJ;
                                                                                                                                                                                                                                               100.0%; llarity 100.0%; Conservative 0;
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                                                                                                                                                                                                                                              Score 27; DB 21; Length 990; Pred. No. 0.04; Indels
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                                                                                                                                                                                                                                                 0;
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   exon
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RESULT 13
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29-OCT-1992;
23-AUG-1994;
                                                15-DEC-1994;
29-OCT-1992;
23-AUG-1994;
12-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This DNA sequence comprises a 992-bp human prostate specific antigen (PSA) cDNA PCR fragment. New Oligonucleotide primers (AAT91044-53) are designed to amplify the PSA gene but not the human glandular kallikrein gene, which has 75-85% homology to the PSA gene. The primers can be used for diagnostis of prostate cancer micrometastasis. A claimed diagnostic kit for detecting prostate cancer comprises a pair of primers, where one of the primers is selected from those given in AAT91044-53, and a means for visualising amplified DNA (ethidium bromide stain, 32p or biotin). The invention also includes a method of detecting cells which express prostate cancer specific sequences by detecting the presence of nucleic acids specific for prostate cancer or a signal specific for
                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                                                     Prostate specific antigen; PSA; D1 stage prostate cancer; D2 stage prostate cancer; D3 stage prostate cancer; human; diagnosis; glandular kallikrein gene; micrometastasis; detection; ss.
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(UYJE-) UNIV
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94US-0294611.
96US-0764527.
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RESULT 14
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                                                                                                                                                                                                                                                                                       dermatological; antiallergic; antiasthmatic; antidiabetic; cytostatic; neuroprotective; antidepressant; nootropic; antiparkinsonian; infection; immunostimulant; gene therapy; antisense therapy; vaccine; inflammation; antianaphylactic; rheumatoid arthritis; septic shock; pancreatitis; cardiac dysfunction; neuropathology; cardiac anaphylaxis; autoimmunity; genetic disease; haematopoietic disorder; platelet disorder; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                           Human; cancer; ulcer; HIV infection; human immunodeficiency virus; antiinflammatory; antirheumatic; antiarthritic; immunosuppressive; antibacterial; endocrine; cardiant; central nervous system; virucide; anti-HIV; fungicide; antimutagen; cardiovascular; antianaemic; anaemia;
                                                                                                                                                                                                                    thrombocyropaenia; osteoporosis; severe combined immunodeficiency; allergic rhinitis; diabetes; multiple sclerosis; depression; Alzheimer's disease; Parkinson's disease; neurodegenerative disord neurological disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                               antiaggregant; haemostatic; vulnerary; antiulcer; osteopathic; eczema;
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23-DEC-1999; 21-JAN-2000; 25-APR-2000;

99US-0471275. 2000US-0488725. 2000US-0552317.

22-DEC-2000;

2000WO-US35017.

2001-457603/49. YT, Liu C, Drmanac

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC AAH99166 to AAH99904 encode the human proteins given in AAM2525 to CC AAM25963. The proteins can have activities based on the tissues and CC cells they are expressed in, such as: antinflammatory; antirheumatic; CC antiarrhritic; immunosuppressive; antipacterial; endocrine; cardiant; central nervous system, virucide; anti-HTV, fungicide; antiaterary; CC antidiabetic; cytostatic; dermatological; antiallergic; antiasthmatic; CC antidiabetic; cytostatic; neuroprotective; antidepressant; noctropic; CC antidiabetic; cytostatic; neuroprotective; antidepressant; noctropic; CC antidiabetic; cytostatic; neuroprotective; antidepressant; noctropic; CC antidiabetic; not munostimulant. *The proteins and polynucleotides cenceding them can be used in gene therapy, antisense therapy and vaccine croding them can be used in gene therapy, antisense therapy and vaccine condinction. The proteins and polynucleotides are useful for screening for agonists or antagonists of a protein and for the treatment and diagnosis of alcorders associated with the activity of a protein e.g. inflammation, constrained arthritis, septic shock, pancreatitis, cardiac dysfunction, confinently, cardiac anaphylaxis, viral, bacterial, HTV and fungal infections, autoimmunity, genetic disease, haematopoietic discorders, canaemia, platelet disorders, thrombocytopaenia, wounds, burns, ulcers, csteeporosis, severe combined immunodeficiency, eczema, allergic CC nlinents, asthma, diabetes, cancer, multiple sclerosis, depression, cCC neuropaical disorders, Parkinson's disease, neurodegenerative and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                               Human; 5' EST; expressed sequence tag; secreted protein; cDNA isolation; gene therapy; chromosome mapping; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 604-605; 1217pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated human polynucleotides encoding treatment and diagnosis of e.g. cancer,
WPI; 2000-500381/45
                                                                                                                                                                                                                EP1033401-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1040 BP; 194 A; 287 C; 302 G; 257 T; 0 other;
                                   Dumas Milne Edwards
                                                                  (GEST ) GENSET
                                                                                                      26-FEB-1999;
                                                                                                                                         21-FEB-2000; 2000EP-0200610
                                                                                                                                                                             06-SEP-2000.
                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                         Human secreted
                                                                                                                                                                                                                                                                                                                                                                          06-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAC03740 standard; cDNA; 1066 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  al Similarity
27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 27; Dilarity 100.0%; Pred. No. 0.0 Conservative 0; Mismatches
                                                                                                          99US-0122487
                                                                                                                                                                                                                                                                                                                                       protein 5' EST, SEQ ID NO: 3738.
                                   Ĺ
                                   Duclert A,
                                   Giordano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptides, useful for ulcers and HIV infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 22; Length 1040;
0.04;
                                   Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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The present sequence is one of a large number of 5′ ESTs derived from mRNAs encoding secreted proteins. An ORF has been identified within the sequence. The 5′ ESTs were prepared from total human RNAs or polyAr RNAs derived from 30 different tissues. EST sequences usually correspond mainly to the 3′ untranslated region (UTR) of the mRNA because they are often obtained from oilgo-dT primed cDNA libraries. Such ESTs are not well suited for isolating cDNA sequences derived from the 5′ ends of mRNAs and even in those cases where longer cDNA sequences have been obtained, the full 5′ UTR is rarely included. 5′ ESTs are derived from mRNAs with intact 5′ ends and can therefore be used to obtain full length cDNAs and genomic DNAs. 5′ ESTs are also used in diagnostic, forensic, gene therapy and chromosome mapping procedures. They are used to obtain vectors.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID 3738; 71pp + CD-ROM; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acid that is a 5' expressed sequence tag (5' EST) for obtaining cDNAs and genomic DNAs that correspond to 5'ESTs and for diagnostic, forensic, gene therapy and chromosome mapping procedures
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P-PSDB; AAG03734
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Ωy Matches Query Match Best Local Sequence 1066 BP; 230 A; 283 C; 310 G; 241 T; 2 other; 1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27 tch 100.0%; Score 27; I al Similarity 100.0%; Pred. No.:0 27; Conservative 0; Mismatches 0; Mismatches DB 21; .04; 0; Length 1066; Indels 0 Gaps

0;

Search completed: November 19, Job time: 193.769 secs 2002, 03:06:38

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В

199 GTTCTGGTGCACCCCCAGTGGGTCCTC 225

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Result
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Database
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Minimum DB seq
Maximum DB seq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OM nucleic - nucleic search, using
                        C
                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                            Score
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Match
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27
1 gttctggtgcacccccagtgggtcctc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                November 19, 2002, 01:43:19; Search time 1542.46 Seconds (without alignments) 283.494 Million cell updates/sec
 100.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IDENTITY_NUC
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em_estov:*
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em_gss_hum:*
em_gss_inv:*
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gb_htc:*
gb_est3:*
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 316
336
371
377
397
435
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 AA593245
AA506459
AA503943
AA603297
AA573604
AA177005
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Compugen Ltd
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AA593245 nn07e10.s
AA506459 nh46a08.s
AA506459 nh46a08.s
AA603297 np12c11.s
AA603297 np12c11.s
AA573604 nf42d12.s
AA177005 nc03b02.s
                                                                        Description
                                                                                                                                         REFERENCE
AUTHORS
TITLE
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VERSION
KEYWORDS
SOURCE
ORGANISM
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AA593245
LOCUS
DEFINITION
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                                                                                                                        JOURNAL
                                                                                                                                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.
Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                    Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 316)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
NATIONAL Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                               AA593245

316 bp mRNA linear EST 25-SEP-1997
nn07e10.s1 NCI_CGAP_Pr4.1 Homo sapiens cDNA clone IMAGE:1077066
similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                                                                                                          Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                mRNA sequence.
AA593245
AA593245.1 GI:2409007
 human.
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AA578976
2 BE840737
2 BE6407537
2 BF679591
AI557591
2 BF679511
AI525089
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BF679168
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AA573575
BE840533
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AI685510
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BF678386
BF678386
BF675112
BF675142
BF679394
BF6781507
BF678076
BF678076
BF67376
BF67376
BF673773
BF6737373
BF673243
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                                                                                                                                                                                                                                                                                                                                              ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                          AA533652 nj72908.s
AI68510 tu36b10.x
BI046475 MR3:PN020
AI525128 promrna-7
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AA573575 nf41f10.s
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BE840533 RC1-FN018
BA639901 np08f03.s
BE840701 RC1-FN018
BE840707 RC1-FN018
BE840707 RC1-FN018
BE767591 pt2.1-1.F
BF679591 pt2.1-1.F
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BF675323 602138273
BF675323 602138273
BF675523 602263138282
BF6755331 602138288
BF6755331 602138288
BF6755331 602138288
                                                                                                                                                                           Euteleostomi;
Homo.
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FEATURES
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AUTHORS
TITLE
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VERSION
KEYWORDS
SOURCE
ORGANISM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 2
AA506459
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ORIGIN
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Best Local :
                                                                                                                                                         source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               source
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tumor Gene Index

Unpublished (1997)

Lamail: cgapbs-remail.nih.gov

Tissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: David B. Krizman, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-GAP clone distribution information can be

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 442 Std Error: 0.00

Seq primer: -40ml3 fvd. ET from Amersham.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cal Similarity 100 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AA506459 mRNA linear EST 20-AUG-1997 nh46a08.sl NCI_CGAP_Pr5 Homo sapiens cDNA clone IMAGE:955382 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 336)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA sequence.
AA506459
AA506459.1 GI:2242699
EST
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h quality sequence stop: 293.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /lab_host="DH10B"
/lab_host="DH10B"
/note="Organ: prostate; Vector: pamp10; mrNA made from /note="Organ: prostate; intraepithelial neoplasia (high-grade), cDNA made by oligo-dT priming. Non-directionally cloned.
Size-selected on agarose gel, average insert size 600 bp.
CDNA Library Preparation: David B. Krizman, ph.D.
Reference: Krizman et al. (1996) Cancer Research, ph.D.
56:5380-5383. cDNA Library Arrayed by: Greg Lennon, ph.D.
DNA Sequencing by: Washington University Genome Sequencing
Center
                 /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:955382"
/clone_lib="NCI_CGAP_Pr5"
/sex="malle"
/tissue_type="prostate"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /tissue_type="prostatic intraepithelial neoplasia - high grade"
                                                                                                                                                                      Location/Qualifiers
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/db_xref="taxon:9606"
/clone="IMAGE:1077066"
/clone_lib="NCI_CGAP_Pr4.1"
/sex="male"
  'note="Vector: pAMP10; mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 27; DB
Pred. No. 0.8
0; Mismatches
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0.89;
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from normal
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AUTHORS
TITLE
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ORGANISM
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AA503943
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QΥ Дb

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Query Match 100.
Best Local Similarity 100.
Matches 27; Conservative
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Tumor Gene Index

Unpublished (1997)

Email: cgapbs-remail.nih.gov

Email: cgapbs-remail.nih.gov

Tissue Procurement: David G. Bostwick, M.D., Rodrigo F. Chuaqui,

M.D., Michael R. Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: David B. Krizman, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Mashington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html
Insert Length: 454 Std Error: 0.00

Seq primer: -40m13 fwd. ET from Amersham

High quality sequence stop: 365.

Std Carlor Qualifiers
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                                                                                                       l Similarity
27; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AA503943 371 bp mRNA linear EST 20-AUG-1997 nh38d05.sl NCI_CGAP_Pr5 Homo sapiens cDNA clone IMAGE:954633 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 371)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
Mational Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                             /tissue_type="prostate"
/lab_host="DH10B"
/lab_host="bH10B"
/lab_host="bH10B"
/note="Vector: pampl0; mRNA made from normal prostatic note="Vector: pampl0; mRNA made by oligo-dT priming.

kon-directionally cloned. Size-selected on agarose gel, average insert size 600 bb. "

a 112 c 111 g 85 t
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Non-directionally colons. Size-selected on agarose gel,
average insert size 600 bp. "
                                                                                                                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:954633"
/clone_lib="NCI_CGAP_Pr5"
/sex="male"
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                                                                                                          0;
                                                                                                       Score 27; DB 9;
Pred. No. 0.9;
; Mismatches 0
                                                                                                          0;
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                                                                                                                                                      Length 371;
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RESULT 5
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DEFINITION
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TITLE
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Best Local Similarity
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                        ORGANISM
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                                                                                                                                                                                                                                                                                                                   GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                                                                                                                                                                                                                             GTTCTGGTGCACCCCCAGTGGGTCCTC 262
                                                                                                                                                                                                                                                                                                                                                                              27;
                                                                                                                                           AA573604 397 bp mRNA linear EST 12-SEP-1997 nf42d12.sl NCI_CGAP_Pr2 Homo sapiens cDNA clone IMAGE:916439 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                                                               EST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    www-bio.llnl.gov/bbrp/image/image.html.
Insert Length: 449 Std Error: 0.00
Seq primer: -40ml3 fwd. ET from Amersham
High quality sequence stop: 345.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AA603297 377 bp mRNA linear EST 21-OCT-1997 np12c11.s1 NCI_CGAP_Pr3 Homo sapiens cDNA clone IMAGE:1116116 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                   Homo sapiens
                                                                                  AA573604.1 GI:2348132
                                                                                                      mRNA sequence.
AA573604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CDNA Library Preparation: David B. Krizman, Ph.D. CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D. DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-GGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D., Michael Emmert-Buck, M.D., Ph.D.
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Unpublished (1997)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                           human.
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National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note-"Vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st /note-"Vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st strand cDNA was primed with oligo(dT)17 on 50 ng of DNAse-treated, total cellular RNA obtained from 5,000-10,000 microdissected cells histologically-determined to be fully malignant prostate cancer cells. Double-stranded cDNA was ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1116116"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /dev_stage="45 years old"
/lab_host="DH10B"
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Pred. No. 0.9;
0; Mismatches
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ORGANISM
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AA177005
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Best Local
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95 GTTCTGGTGCACCCCCAGTGGGTCCTC 121
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                                                                                                  Homo sapiens

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 435)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA177005 435 bp mRNA linear EST 14-AUG-19: nc03b02.s1 NCI_CGAP_Pr2 Homo sapiens cDNA clone IMAGE:302 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);, mRNA
Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
                                                            Unpublished (1997)
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High quality sequence stop: 350.
Location/Qualifiers
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Tissue Procurement: W. Marston Linehan, M.D.,
Michael Emmert-Buck, M.D., Ph.D.
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Unpublished (1997)
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Insert Length: 1526 Std Error: 0.00
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/clone_lib="NCI_CGAP_Pr2"
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/db_xref="taxon:9606"
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Pred. No. 0.
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0.9;
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Marston Linehan, M.D., Rodrigo Chuaqui, M.D.

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KEYWORDS
SOURCE
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AUTHORS
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AA533652
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                                                                                                                                         Contact: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo F. Chuaqui,
M.D., Michael R. Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Catarrhini; Hominidae; Homo. 1 (bases 1 to 453) 1 (bases 1 to 453) NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap. NATIONAL Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Michael Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: David B. Krizman, Ph.D.

cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.

cDNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.ilni.gov/bbrp/image/image.html

Seq primer: 40M13 fwd. from Amersham.

Location/Qualifiers
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A53 bp mRNA linear EST 21-AUG-1997
n)72g08.sl NCI_CGAP_Pr10 Homo sapiens cDNA clone IMAGE:998078
similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
27; Conser
Trace considered overall poor quality
Insert Length: 1654 Std Error: 0.00
Seg primer: -40ml3 fwd. ET from Amersham
                                                                                                                                                                                                                                                                                                                                                                                                       Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA sequence
AA533652
                                                                                                               www-bio.llnl.gov/bbrp/image/image.html
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64 a 119 c 146 g 105 t 1 others
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/db_xref="taxon:9606"
/clone="IMAGE:302"
/clone_lib="NCI_CGAP_Pr2"
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/lab_host="DH10B"
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VERSION
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AI685510/c
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Emmail: cgapbs-r@mail.nih.gov

Tissue Procurement: Michael J. Brownstein, M.D., Ph.D., Michael R.

Emmert-Buck, M.D., Ph.D.

CDNA Library Preparation: M. Bento Soares, Ph.D.

CDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Seq primer: -400P from Gibco.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 454)
1 (bases 1 to 454)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mRNA sequence.
AI685510
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Location/Qualifiers
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Unpublished (1997)
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/note-"Organ: prostate; vector: pf773D-Pac (Pharmacia) with a modified polylinker; Plasmid DNA from the normalized library NCI_CGAP_P722 was prepared, and sscircles were made in vitro. Following HAP purification, this DNA was used as tracer in a subtractive hybridization reaction. The driver was PCR-amplified cDNAs from a pool of 5,000 clones made from the same library (cloneIDs 985608-986759, 1101192-1101959, and 1217928-1220615).
                                                                                                                                                                                                                                                                                               /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:2253115"
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                                                                                                                                                                                                     /dev_stage="adult"
/lab_host="DH10B"
                                                                                                                                                                                                                                                  /clone_lib="NCI_CGAP_Pr28"
/sex="male"
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/db_xref="taxon:9606"
/clone="IMAGE:998078"
/clone_lib="NCI_CGAP_Pr10"
/sex="male"
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/lab_host="DH10B"
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                     1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
      GTTCTGGTGCACCCCCAGTGGGTCCTC 349
                                                                                                                                                                                                                                                                                                                                                                                              Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pi?tl=MR3&t2-MR3-FN0209-
070201-010-b11&t3=2001-02-07&t4=1)
Seq primer: puc 18 forward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dias Neto,E., García Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagal,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HR3-FN0209-070201-010-b11 FN0209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Simpson, A.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence tags
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (bases
                                                                                                                                                                                                                                                                                                                                              quality sequence start: 7 quality sequence stop: 465. Location/Qualifiers
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                                                                                                                                              83
                                                                   Conservative
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                                                                                                                                           /note="Organ: prostate_normal; Vector: puc18; Site_1: SmaI; Site_2: SmaI; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

134 c 144 g 103 t 1 others
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                                                                                                                                                                                                                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="FN0209"
                                                                                                                                                                                                                                                                        /dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 to 465)
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                                                                                100.0%;
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AUTHORS
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SOURCE
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LOCUS
DEFINITION
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KEYWORDS
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui,
, Michael Emmert-Buck, M.D., Ph.D.
cDNA Library Preparation: David B. Krizman, Ph.D.
cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.
                                                                                Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                          Mammalia; Eutheria; Primates;
1 (bases 1 to 484)
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                                                                                                                                                                                                                                                         AA603529.1
                                                                                                                                                                                                                                                                        mRNA sequence
AA603529
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Fax: 5106280108
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Mammalia; Eutheria;
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AI525128
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                                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Leroy Hood
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University of Washington
Department of Molecular Biotechnology,
Washington, Seattle, WA 98195
                                                                                                                                                                                                                          AA603529 484 bp mRNA linear EST 08-OCT-1997 np15b08.sl NCI_CGAP_Pr3 Homo sapiens cDNA clone IMAGE:1116375 similar to gb:M21895 PROSTATE SPECIFIC ANTIGEN PRECURSOR (HUMAN);,
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Prostate cancer expression profiling Genomics 59 (2), 178-186 (1999)
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Huang,G.M., Ng,W.l., Farkas,J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="bvtumor"
/clone_lib="bvtumor"
/note="Organ: Prostate; Vector: pBluescript; DcDNA library was constructed using Lambda ZP I (Stratagene). mRNA was extracted from a fresh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tumor tissues (Urology Department, University of Washington) "
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/db_xref="taxon:9606"
                                                           Chordata; Craniata; Vertebrata; Primates; Catarrhini; Hominidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                             Hominidae;
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  Project (CGAP),
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                                                                                Euteleostomi;
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Ph.D.

M. D.

source

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Tumor Gene Index

Unpublished (1997)

Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D.

Michael Emmert-Buck, M.D.; Ph.D.

CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Preparation: David B. Krizman, Ph.D.

CDNA Library Arrayed by: Genome Systems Inc., Greg Lennon, Ph.D.

CDNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be

Tound through the I.M.A.G.E. Consortium/LLNL at:

WWW-bio.llnl.gov/bbrp/image/image.html
                                Trace considered overall poor quality Insert Length: 1535 Std Error: 0.00 Seq primer: -40ml3 fwd. Er from Amersham High quality sequence stop: 1.

Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (Dases 1 to 485)

1 (NGI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

NGI-CGAP institute, Cancer Genome Anatomy Project (CGAP),
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AA573575
AA573575.1 GI:2348103
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AA573575
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ANTIGEN PRECURSOR (HUMAN);
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/db_xref="taxon:9606"
/clone="IMAGE:1116375"
/clone_lib="NCI_CGAP_Pr3"
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/lab_host="DH10B"
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/sex="Male"
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Pred. No. 0.91;
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AUTHORS
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: +55-11-2707001

Tel: +55-11-2707001

Ten: +55-11-2707001

Email: asimpson@ludwig.org.br

This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?ti=st2=RC1-FN0188-260 700-022-f06&t3=2000-07-26&t4=1)

Seq primer: puc l8 forward High quality sequence stop: 495.

Location/Qualiflers

1.495
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20202663
Contact: Simpson A.J.G.
Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Ludwig Institute For Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 495)

Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R., Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F., Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H., Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     al Similarity 100
27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence tags
Proc. Natl. Acad. Sci. U.S.A. 97 (7),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Simpson,A.J.
Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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/organism="Homo sapiens"
/olone="ImAge:91363"
/clone="ImAge:91363"
/clone=lib="NCI_CGAP_Pr2"
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NCI_CGAP_Pr2"
/sex="Male"
/sex=
/organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="FN0188"
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tive 0;
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Pred. No. 0.91;
Mismatches 0
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0

FEATURES

REFERENCE AUTHORS TITLE

ACCESSION VERSION KEYWORDS SOURCE ORGANISM

Homo sapiens

RESULT 12 AA573575 LOCUS DEFINITION

187

ORIGIN BASE COUNT

89

Query Match Best Local S Matches 27

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RESULT 14
BE840441/c
BASE COUNT
ORIGIN
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?t1=&t2=RC1-FN0188-210
700-021-a02&t3=2000-07-21&t4=1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             l (bases 1 to 505)
Dias Neto,E., Garcia Correa,R., Verjovski-Almeida,S., Briones,M.R.,
Nagai,M.A., da Silva,W. Jr., Zago,M.A., Bordin,S., Costa,F.F.,
Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H.,
Brunstein,A., deOliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BE840441
RC1-FN0188-210700-021-a02 FN0188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Shotgun sequencing of the human transcriptome with ORF expressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human.
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+55-11-2707001
                                                                                                                                                                                                                                                                                                                                                             quality
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Natl. Acad. Sci. U.S.A. 97 (7), 3491-3496 (2000)
               application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions."

164 c 136 g 78 t
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a 161 c 131 g 77 t 1 others
                                                                                                                                        /note="Organ: prostate_normal; Vector: puc18; Site_1: SmaI
; Site_2: SmaI; A mini-library was made by cloning
products derived from ORESTES PCR (U.S. Letters Patent
                                                                                                                                                                                                                                        /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="FN0188"
                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                  /dev_stage="Adult"
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Query Match

Score 27;

DВ

12;

Length 505;

Search completed: Job time: 1545.46

November 6 secs

19,

2002, 05:17:31

В Q

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SOURCE
ORGANISM
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  187
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                                                                                                   Local
                 1 GTTCTGGTGCACCCCCAGTGGGTCCTC
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                                                                               27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          www-bio.llnl.gov/bbrp/image/image.html
Seq primer: -40m13 fwd. ET from Amersham
High quality sequence stop: 367.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Robert Strausberg, Ph.D.
Email: Gapbs-remail.nih.gov
Tissue Procurement: W. Marston Linehan, M.D., Rodrigo Chuaqui, M.D., Michael Emmert-Buck, M.D., Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tumor Gene Index Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA639901 523 bp mRNA linear EST 23-OCT-199 np08f03.sl NCI_CGAP_Pr3 Homo sapiens cDNA clone IMAGE:1115741 similar to gb:m21895 prostate SPECIFIC ANTIGEN PRECURSOR (HUMAN);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalia; Eutheria;
1 (bases 1 to 523)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA Library Preparation: David B. Krizman, Ph.D. cDNA Library Arrayed by: Genome Systems Inc., Greg Lennon,
                                                                                                                                                                                     100
                                                                                  Conservative
                                                                                                                                                                          /note="vector: pAMP10; Site_1: Not1; Site_2: EcoRI; 1st strand cDNA was primed with oligo(dT)17 on 50 ng of DNAse-treated, total cellular RNA obtained from 5,000-10,000 microdissected cells histologically-determined to be fully malignant prostate cancer cells. Double-Stranded cDNA was ligated to EcoRI adaptors, 5 cycles of PCR applied to the cDNA with an adaptor-specific primer, and the resulting PCR product subcloned into pAMP10 by the UDG-cloning method (Life Technologies). Average insert size is 600 bp. NOTE: Not directionally cloned. This library was constructed by David Krizman."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:1115741"
/clone_lib="NCI_CGAP_Pr3"
                                                                                                                                                                                                                                                                                                                                                                                                                                  /dev_stage="45 years old"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /sex="Male"
                                                                                                   100.0%;
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                                                                                  Mismatches
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                                                                                                                        Length 523;
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                                                                               Gaps
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Minimum
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No.
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Maximum Match 100%
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Perfect score:
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                                                Issued_Patents_NA:*
1: /cgn2_6/ptodata/2,
2: /cgn2_6/ptodata/2,
3: /cgn2_6/ptodata/2,
4: /cgn2_6/ptodata/2,
5: /cgn2_6/ptodata/2,
6: /cgn2_6/ptodata/2,
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length: 2000000000
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Copyright (c) 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gttctggtgcaccccagtgggtcctc
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/BCTUS_COMB.seq:*
/cgn2_6/ptodata/2/ina/backfiles1.seq:*
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          US-08-931-981A-1
US-09-100-264-4
US-08-358-782D-14
US-08-358-782D-13
US-08-605-785-616
US-08-605-785-616
US-08-605-785-105-31
US-08-61-527-0468-2
US-08-622-0468-2
US-08-622-0468-13
US-09-100-264-2
US-08-767-820A-7
PCT-US-5-6157-7
US-08-622-0468-17
US-08-62-0468-17
US-08-767-820A-20
US-08-767-820A-20
US-08-767-820A-20
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Sequence 1, Appli
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Sequence 14, Appl
Sequence 14, Appli
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120 GTTCTGGTGCACCCCCAGTGGGTCCTC

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QY 1 GTTCTGGTGCACCCCAGTGGGTCCTC 27	Query Match 100.0%; Score 27; DB 1; Lengest Local Similarity 100.0%; Pred. No. 0.0062; Matches 27; Conservative 0; Mismatches 0; I	INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 200 base pairs TYPE: nucleic acid STRANDEDNESS: double TOPOLOGY: linear MOLECULE TYPE: genomic DNA (PSA exons 2 and 3) US-08-931-981A-1	CLASSIFICATION: 536 ATTORNEY/AGENT INFORMATION: NAME: Paul D. Yasger REGISTRATION NUMBER: 37,477 REFERENCE/DOCKET NUMBER: 6179.US.O1 TELECOMMUNICATION INFORMATION: TELEPHONE: 847/938-3508 TELEPAX: 847/938-2623	ZIP: 60064-3500 COMPUTER READABLE FORM: MEDJUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION UNBER: US/08/931,981A FILING DATE:	RESULT 1 US-08-931-981A-1 US-08-931-981A-1 Sequence 1, Application US/08931981A Patent No. 5817798 GENERAL INFORMATION: APPLICANT: 6. Gundling TITLE OF INVENTION: RAPID RNA ISOLATION PROCEDURE NUMBER OF SEQUENCES: 4 CORRESPONDENCE ADDRESS: ADDRESSEE: Abbott Laboratories STREET: 100 Abbott Park Road CITY: Abbott Park STATE: Illinois COUNTRY: USA	19.6 72.6 234 4 19.6 72.6 234 4 19.6 72.6 700 4 19.6 72.6 765 4 19.6 72.6 765 4 19.6 72.6 871 1	28 23.8 88.1 832 3 US-08-622-046B-4 29 23.8 88.1 832 3 US-08-622-046B-15 30 23.8 88.1 832 5 PCT-US95-06157-5 31 23.8 88.1 1341 4 US-08-983-075D-6 32 23.8 88.1 1358 4 US-08-983-075D-8 32 23.8 88.1 1358 4 US-08-983-075D-8 33 21 77.8 21 4 US-09-493-491-11 34 21 77.8 21 4 US-09-493-491-11 35 20.6 73.3 1364 4 US-09-280-116-20 36 19.6 72.6 45 4 US-09-280-116-20 37 19.6 72.6 234 4 US-09-605-785-328 38 19.6 72.6 234 4 US-09-605-785-328 39 19.6 72.6 234 4 US-09-605-785-328
	Length 200; Indels 0; Gaps C					2, 2, 2, 3, 3, 3, 3, 3, 3, 3, 3, 3, 3, 3, 3, 3,	Sequence 4, Appli Sequence 15, Appli Sequence 5, Appli Sequence 6, Appli Sequence 8, Appli Sequence 11, Appli Sequence 24, Appli Sequence 20, Appli Sequence 790, Appli Sequence 328, Appli Sequence 328, Appli
	0;		•				

RESULT 2 US-09-100-264-4

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APPLICANT: Young, Charles Y.F.
APPLICANT: Tindall, Donald J.
APPLICANT: Klee, George G.
TITLE OF INCENTION: Method for detection of breast cancer
TITLE REFERENCE: 545.006US1
CURRENT APPLICATION NUMBER: US/09/100.264
CURRENT FILING DATE: 1998-06-19
EARLIER FAILIG DATE: 1997-06-20
UNMBER OF SEQ ID NOS: 15
SOPTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 4
LENGTH: 711
TYPE: NAME

LENGTH: 711
                                                                           ZIP: 19103
ZIP: 19103
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBN PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING NETE: 15-DEC-1994
FILING DATE: 15-DEC-1994
FLILING DATE: 15-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y
NAME: Beardell, Lori Y
REGISTRATION NUMBER: 34,293
REGISTRATION INFORMATION:
TELEPONE: 215-568-3100
TELEPAN: 215-568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: DNA; Homo sapiens; ORGANISM: Homo sapiens US-09-100-264-4
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Patent No. 6235486
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CHART NO. 5674682
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-358-782D-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis of Prostate
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. Grant
APPLICANT: Moreno, Jose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: One Liberty Place 45th. Floor
CITY: Philadelphia
STATE: PA
ZIP: 'PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85 GTTCTGGTGCACCCCAGTGGGTCCTC 111
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Local Similarity 100.0%; Pred. No. 0.
Les 27; Conservative 0; Mismatches
                           TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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DNA (genomic)
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                                                                                                                                             MOLECULE TYPE: DN
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-764-527A-13
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US-08-764-527A-13
                                                                                                                                                                                                                                                                  PRIOR APPLICATION NOME:
APPLICATION NUMBER: US/08/358,782
FILING DATE: 15-DEC-1994
ATTORNEY/AGENT INFORMATION:
AMME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-1327
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
TELEPAX: 215-568-310
TELEPAX: 215-568-310
TELEPAX: 215-568-310
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 992 base pairs
TYPE: DICIPIC STATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/764,527A
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.
Best Local Similarity 100.
Matches 27; Conservative
                                                                     Query Match
Best Local Similarity
Matches 27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & STREET: One Liberty Place 46th. Floor CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Methods of Detecting Micrometastasis TITLE OF INVENTION: Prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 992 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                       100.0%; Score 27; I ilarity 100.0%; Pred. No. 0 Conservative 0; Mismatches
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100.0%; Pred. No. 0.
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RESULT 5 US-09-605-785-616; A; Sequence 616; A;

Application US/09605785

PPLICANT: Xu, Jiangchun

INFORMATION:

PPLICANT:

Mitcham, Jennifer L. Dillon, Davin C.

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; SOFTWARE: FastSEQ for Windows Version 3.0; SEQ ID NO 616; LENGTH: 1350; TYPE: DNA; ORGANISM: Homo sapien US-09-605-785-616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 6
US-08-358-782D-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Croce, Carlo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/605,785
CURRENT FILING DATE: 2000-06-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: SKEİKY, YASİR A.W.
APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 835
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              ATTORNEY/AGENT INFORMATION:
NAME: BEARDELI, LOTI Y.
REGISTRATION NUMBER: 34, 293
REFERENCE/DOCKET NUMBER: TJU
                                                                              SOFTWARE: PatentIn Release #1.0, VGCURRENT APPLICATION DATA:
APPLICATION UNMERR: US/08/358,782D
FILING DATE: 15-DEC-1994
CLASSIFICATION: 435
                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                            APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis of Prostate
NUMBER OF SEQUENCES: 14
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                           Cli.
STATE: F.,
19103
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nes 27; Conserv
                                                                                                                                                                                                                                                                        CITY: Philadelphia
                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1, Application US/08358782D 5674682
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Vedvick, Thomas S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kalos, Michael D. Fanger, Gary R. Retter, Marc W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wang,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Carter, Darrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stolk, John A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jiang, Yuqui
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Harlocker,
                                                                                                                                                                                                                                                                                           E: Woodcock Washburn Kurtz Mackiewicz & One Liberty Place 46th, Floor
                                                                                                                                                                                                                                                                                                                                                                                              Moreno, Jose
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            TJU-1327
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                                                                                                                                                    Version #1.25
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0.0072;
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; HYPOTHETICAL:
; ANTI-SENSE: NO
US-08-764-527A-14
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Best Local Similarity
Matches 27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                   TELEFAX: 215-568-3439 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
                                                                                                                                                                                                REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJ
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-3100
                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08
FILING DATE: 15-DEC-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                          MOLECULE TYPE:
                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cli.
STATE: **.
19103
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                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz STREET: One Liberty Place 46th. Floor
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                                                                                            TOPOLOGY:
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                                                                                                                                       LENGTH:
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Length 1462;
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Query Match Best Local Similarity

100.0%;

Score Pred.

27; No.

DB 2; 0.0073;

В QY

Matches

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; NAME/KEY:
; LOCATION:
US-08-844-024-1
                                                                                RESULT 9
US-08-718-547-1
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION UNBER: 08/229,391
EILING DATE: 15-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 43677/J
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEFAX: (212) 664-0525
TELEX: 42253 COOP UI
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1729 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/08844024 Patent No. 5840494
                             Sequence 1, Application US/08718547 Patent No. 5976794 GENERAL INFORMATION:
                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LIBERT: 30 Rockefeller Plaza
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10112
MPUTER PT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Katz, Aaron E., et al.
TITLE OF INVENTION: A Method For Molecular Staging Of
TITLE OF INVENTION: Prostate Cancer
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: lin
MOLECULE TYPE:
ORIGINAL SOURCE:
APPLICANT: Katz, Aaron E.; Buttyan, Ralph; Raffo, Anthony; APPLICANT: Olsson, Carl A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              181
                                                                                                                                             APPLICATION NUMBER: US/08/844,024 FILING DATE: 18-APR-1997
                                                                                                                                                                                                                                                                                                                                             ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: S1
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
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378..1088
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Pred. No. 0.0
D; Mismatches
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                                                             APPLICANT: KUPS, THOMAS J.

APPLICANT: WU, YUNQI

TITLE OF INVENTION: VACCINES WITH ENHANCED INTRACELLULAR

TITLE OF INVENTION: PROCESSING

FILE REFERENCE: 233/221

CURRENT APPLICATION NUMBER: US/09/056,105

CURRENT FILING DATE: 1998-04-06

EARLIER APPLICATION NUMBER: 60/043,467

EARLIER FILING DATE: 1997-04-10

NUMBER OF SEQ ID NOS: 35

SOFTWARE: FastSEQ for Windows Version 3.0.

SEQ ID NO 31

LENGTH: 7130

TYPE: DNA

CENTIFY: 130
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; LOCATION:
US-08-718-547-1
                                   ; ORGANISM: Homo sapiens US-09-056-105-31
                                                                                                                                                                                                                                                                                                                                             RESULT 10
US-09-056-105-31
                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                             Sequence 31, Application US/09056105 Patent No. 6287569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local
   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX:
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENCTH: 1729 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 43
REFERENCE/TOOKET NUMBER: 413
TELECOMMUNICATION INFORMATION
TELEPHONE: (212) 278 0400
TELEFAX: (212) 391-0525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIE: 10036

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/718,547
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 1185 Avenue of the Americas CITY: New York
STATE: New York
COUNTRY: U.S a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LECULE TYPE:
IGINAL SOURCE:
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27; Conserv
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PE: cDNA
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     Score 27;
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Pred. No. 0.
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0.0074;
es 0;
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     Length 7130;
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Best Local Similarity Matches 27; Conserv

Conservative

100.0%; Pred. No. 0.0082; No. 0.0082;

Indels

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US-08-622-046B-13
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US-08-622-046B-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/08622046B Patent No. 6103237
                                                                                                                                                      Sequence 13, Application US/08622046B Patent No. 6103237
                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 612-339-3061
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Stable Variant NUMBER OF SEQUENCES: 21
                                                                                                         APPLICANT: Saedi, Mohamma APPLICANT: Mikolajczyk,
                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 711 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 612-373-6903
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Saedi, Mohamma APPLICANT: Mikolajczyk,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Stable Variant HK2 Polypeptide
                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 11 MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: FILING DATE: 26-MAR CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                            85 GTCCTGGTGCACCCCCAGTGGGTGCTC 111
                                                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Woessner, Warren D.
REGISTRATION NUMBER: 30,440
REFERENCE/DOCKET NUMBER: 476.001US1
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                            CITY:
                                         STREET:
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25; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                      linear
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                                         Schwegman,
O. Box 2938
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92.6%;
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                                                     Lundberg,
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                                                                                                                                                                                                                                                                                                Score 23.8; DB Pred. No. 0.15; 0; Mismatches
                                                                                                                                                                                                                                                                                                  0;
                                                       Woessner & Kluth, P.A.
                                                                                               HK2 Polypeptide
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Query Match
Best Local Similarity
Matches 25; Conserva
RESULT 14
US-08-768-859A-7
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                                                                                                                                                                                        ; TYPE: DNA; Homo sapiens US-09-100-264-2
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US-09-100-264-2
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                                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 2 LENGTH: 711
                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Young, Charles Y.F.
APPLICANT: Tindall, Donald J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/09100264 Patent No. 6235486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/100,264
CURRENT FILING DATE: 1998-06-19
EARLIER APPLICATION NUMBER: US 60/050,963
EARLIER FILING DATE: 1997-06-20
                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Method for detection of FILE REFERENCE: 545.006US1
                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 15
                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Klee,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 612-339-3061 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: cDNA
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LENGTH: 711 base pair:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Woessner, Warren D.
REGISTRATION NUMBER: 30,440
REFERENCE/DOCKET NUMBER: 476.001US1
                                                               85 GTCCTGGTGCACCCCCAGTGGGTGCTC 111
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EDNESS: double
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92.6%;
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Pred. No. 0.
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Sequence 7, Application US/08768859A Patent No. 6013471
GENERAL INFORMATION:

APPLICANT:

Tindall, Donald J. Young, Charles Y.F.

```
STATE.

COUNTRY: USA
ZIP: 55402

ZIP: 55402

COMPUTER READABLE FORM:

MEDJIM TYPE: Floppy disk

COMPUTER: IBM PC COMPATION

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PASIENTIA Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/768,859A

FILING DATE: 17-DDECHMER-1995

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

REFERENCE/DOCKET NUMBER: 30,440

REFERENCE/DOCKET NUMBER: 355.002US3

TELECOMMUNICATION INFORMATION:

TELEPHONE: 612-339-3061

TELEPHONE: 612-339-3061

TELEPHONE: 612-339-3061

TELEPHONE: 612-339-3061

TELEPHONE: 612-339-3061
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; LOCATION: 7..720
US-08-768-859A-7
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US-08-767-820A-7
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INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 760 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOT.ECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 7, Application US/08767820A Patent No. 6093796 GENERAL INFORMATION:
STATE: MN
COUNTRY: USA
COUNTRY: USA
ZIP: 55402
ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/767,820A
FILING DATE: 17-DECEMBER-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                           APPLICANT: Tindall, Donald J.
APPLICANT: Young, Charles Y.F.
APPLICANT: Saedi, Mohammed S.
TITLE OF INVENTION: DNA Encoding HK2 Variant Polypeptide:
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS: Schwegman, Lundberg, Woessner & Kluth, P.A.
STREET: P.O. Box 2938
CITY: Minneapolis
STATE: MI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Saedi, Mohammed S.
TITLE OF INVENTION: DNA Encoding HK2 Variant Polypeptides
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: P.O. Box :
CITY: Minneapolis
STATE: MN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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25; Conservative
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                                                                                                                                                                                                                                                                                                                          DNA Encoding HK2 Variant Polypeptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            88.1%; Score 23.8; DB 3; Length 760; 92.6%; Pred. No. 0.15; tive 0; Mismatches 2; Indels
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В
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; LOCATION: 7..720
US-08-767-820A-7
                                             Query Match

Best Local Similarity 92.6

Matches 25; Conservative
                                                                                                                                                                                                                                                                      TELEPHONE: 612-339-0331
TELEPAX: 612-339-3061
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:

NAME: WOOSSNEY, WAITEN D.

REGISTRATION NUMBER: 30,440

REFERENCE/DOCKET NUMBER: 545.002US2

TELECOMPUNICATION INFORMATION:

TELEPHONE: 612-339.0331
                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: cDNA
                                                                                                                                                                              FEATURE:
94 GTCCTGGTGCACCCCCAGTGGGTGCTC 120
                   1 GTTCTGGTGCACCCCCAGTGGGTCCTC 27
                                                                             92.6%;
                                                            Score 23.8; DB 3; Pred. No. 0.15; 0; Mismatches 2;
                                                                                             Length 760;
                                                                Indels
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Search completed: November 19, 2002, 05:20:21 Job time: 53.9615 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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1: /cgn2_6/ptodata/1/pubpna/US

2: /cgn2_6/ptodata/1/pubpna/US

3: /cgn2_6/ptodata/1/pubpna/US

4: /cgn2_6/ptodata/1/pubpna/US

5: /cgn2_6/ptodata/1/pubpna/US

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6: /cgn2_6/ptodata/1/pubpna/US

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9: /cgn2_6/ptodata/1/pubpna/US

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13: /cgn2_6/ptodata/1/pubpna/US

14: /cgn2_6/ptodata/1/pubpna/US
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27
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Copyright (c) 1993 - 2002 Compu
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/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US08_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US09_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US10_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq:*
/cgn2_6/ptodata/1/pubpna/US07_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/PCTUS_PUBCOMB.seq:*
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                                                                                                                                                                                                                                                                                                                                   Length
                                                      1350
1350
1350
1729
1758
2406
3426
4661
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861
871
316
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          US-09-829-004A-8
US-09-789-143-616
US-09-780-669-616
US-09-822-827-616
US-09-822-827-708-598
US-09-822-827-980
US-09-755-100-5
US-09-755-100-1
US-09-755-100-1
US-09-755-100-1
US-09-755-300-153
US-09-755-300-153
US-09-925-300-153
US-09-925-300-814
US-09-925-300-814
US-09-822-827-790
US-09-822-827-790
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111.898 Million cell updates/sec
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Sequence 8, Appli
Sequence 616, App
Sequence 616, App
Sequence 977, App
Sequence 980, App
Sequence 980, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 2, Appli
Sequence 153, App
Sequence 153, App
Sequence 101, App
Sequence 207, App
Sequence 790, App
Sequence 790, App
Sequence 790, App
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1248	1248	1215	1215	1215	1214	1214	1214	1214	1167	1167	1167	1167	1167	1119	1119	1119	1119	1119	871	765	765	765	234	234	234
10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10	10
-780-669	US-09-759-143-171	822-827	669	759-143	-115-453	-822-827	-09-780-669	-759-14	US-09-115-453-175	US-09-822-827-175		ċ	US-09-759-143-175	US-09-115-453-177	US-09-822-827-177	us-09-030-606-177	US-09-780-669-177	US-09-759-143-177	US-09-956-999-7	US-09-822-827-524	US-09-780-669-524	US-09-759-143-524	-822-827	780-	US-09-759-143-328
Sequence 171, App		326,	326,	326,	Sequence 225, App	225,	225,	225,	175,	175,	175,	175,	175,	177,	177,	177,	177,	17	7,	52	52	52	32	Sequence 328, App	Sequence 328, App

ALIGNMENTS

RESULT 1 US-09-829-004A-8

Sequence 8, Application US/09829004A Patent No. US20020132976A1 GENERAL INFORMATION:

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Query Match
Best Local Similarity
Thes 27; Conserve
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US-09-759-143-616
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               Sequence 616, Application US/09759143
Patent No. US20020022248A1
GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
APPLICANT: Dillon, Davin C.
                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/829,004A CURRENT FILING DATE: 2001-04-16
PRIOR APPLICATION NUMBER: US 60/195,456
PRIOR FILING DATE: 2000-04-10
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PACENTIN VERSION 3.0
SEQ ID NO 8
LENGTH: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Chong, Pele
APPLICANT: Pedyczak, Artur
APPLICANT: Sia, Charles Dwo Yuan
TITLE OF INVENTION: Immunogenic Peptides Derived from Prostate-Specific Antigen
TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 11014-24
     APPLICANT:
                                                                                                                                                                                                                                                                                                                                        LENGTH: 27
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                    Xu, Jiangchun
Dillon, Davin (
Mitcham, Jenni:
                                                                                                                                                                                                                                   100.0%; ilarity 100.0%; Conservative 0
   Jennifer L.
                                                                                                                                                                                                                                      0;
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Harlocker, Henderson,

Susan L. Robert A. ₽.

Yuqui

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APPLICANT: Hepler, William

APPLICANT: Hural, John

APPLICANT: Howelil, Fatricia D.

APPLICANT: Houghton, Raymond L.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REFERENCE: 210121.427C24

CURRENT APPLICATION NUMBER: US/09/780,669

CURRENT FILING DATE: 2001-02-09

NUMBER OF SEQ ID NOS: 943

SOFTWARE: FastSEQ for Windows Version.3.0

SEQ ID NO 616

LENGTH: 1350

TYPE: DNA

ORGANISM: Homo sapien

US-09-780-669-616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Day, Craig H.

APPLICANT: Vedvick, Thomas S.

APPLICANT: Vedvick, Thomas S.

APPLICANT: Carter, Darrick

APPLICANT: Li, Samuel

APPLICANT: Wang, Aljun

APPLICANT: Weary, Aljun

APPLICANT: Hepler, William

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

FILE REFERENCE: 210121.427023

CURRENT APPLICATION NUMBER: US/99/759,143

CURRENT APPLICATION NUMBER: US/99/759,143

CURRENT APPLICATION NUMBER: US/99/759,143

CURRENT APPLICATION NUMBER: US/99/759,143

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 616

SEQ ID NO 616

LENGTH: 1350

TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 3
US-09-780-669-616
; Sequence 616, Application US/09780669
· Datent No. US20020051977A1
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Best Local S
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APPLICANT:
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APPLICANT:
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APPLICANT:
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Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
Day, Craig H.
Vedvick, Thomas S.
Carter, Darrick
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Fanger, Gary R.
Retter, Marc W.
Stolk, John A.
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Hepler, William
Hural, John
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Vedvick, Thomas S.
Carter, Darrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Harlocker, Susan L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dillon, Davin C.
Mitcham, Jennifer L.
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Pred. No. 0.0087;
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RESULT 4

US-09-822-827-616

US-09-822-827

Sequence 616, Application US/09822827

Patent No. US2002081680A1

GENERAL INFORMATION:

APPLICANT: Xu, Jiangchun

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE FERENCE: 210121.534C1

CURRENT APPLICATION NUMBER: US/09/822,827

CURRENT FILING DATE: 2001-03-28

NUMBER OF SEQ ID NOS: 982

SOFTWARE: FastSEQ for Windows Version 3.0

SERO ID NO 616

LENGTH: 1350
                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/969,708
CURRENT FILING DATE: 2001-10-03
PRIOR APPLICATION NUMBER: US/60/237,606
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: US/60/237,425
PRIOR APPLICATION NUMBER: US/60/237,425
PRIOR FILING DATE: 2000-10-03
NUMBER OF SEQ ID NOS: 658
SOFTWARE: Patentin version 3.0
SEQ ID NO 598
LENGTH: 1729
TYPE: DNA
ORGANISM: Homosapiens
US-09-969-708-598
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                                                                                                                                                                                                                                                                                                                                                                                                Sequence 598, Application US/09969708
Patent No. US20020102532A1
GENERAL INFORMATION:
APPLICANT: Augustus, Meena
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign
TITLE OF INVENTION: Sets
TITLE REFERENCE: 689299-70
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US-09-969-708-598
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Matches 27
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Best Local S
Matches 27
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Best Local (
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778 GTTCTGGTGCACCCCCAGTGGGTCCTC 804
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27; Conserv
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27; Conser
                                                                   100.0%; Score 27; DB 10; Length 1729;
larity 100.0%; Pred. No. 0.0089;
Conservative 0; Mismatches 0; Indels 0
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100.0%; Pred. No. 0
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Pred. No. 0
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488
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0.0087;
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APPLICANT: Xu, Jiangchun
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C1
CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT FILING DATE: 2001-03-28
NUMBER OF SEQ ID NOS: 982
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 977
LENGTH: 1758
TYPE: DNA
ORGANISM: Homo sapiens
US-09-822-827-977
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; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 980
; LENGTH: 2406
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-822-827-980
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US-09-822-827-980
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US-09-755-100-5
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PRIOR APPLICATION NUMBER: IL 128587
PRIOR FILING DATE: 1999-02-18
PRIOR PPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Xu, Jian
                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: SAVITZKY, Kinneret et al.
APPLICANT: SAVITZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P
CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 980, Application US/09822827 Patent No. US20020081680A1
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                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/09755100 Patent No. US2002009189A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.534C1
CURRENT APPLICATION NUMBER: US/09/822,827
CURRENT FILING DATE: 2001-03-28
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US-09-755-100-1
                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/755,100
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: IL 128587
PRIOR FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 131363
PRIOR APPLICATION NUMBER: IL 131363
PRIOR FILING DATE: 1999-08-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 10
US-09-755-100-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 9
US-09-755-100-3
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LENGTH: 3423
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/09755100 Patent No. US20020099189A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 27; DB 10; Best Local Similarity 100.0%; Pred. No. 0.0094; Matches 27; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver. SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: SAVITZKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156p
CURRENT APPLICATION NUMBER: US/09/755,100:
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: IL 128587
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-04-14
PRIOR APPLICATION NUMBER: IL 129439
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-04-14
PRIOR FILING DATE: 1999-08-11
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                                                                                          SOFTWARE:
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                      APPLICANT: SAVITZKY, Kinneret et al. TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES FILE REFERENCE: 2786-0156P
                                                                                                                                         NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQ ID NOS: 12
                   TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1658 GTTCTGGTGCACCCCCAGTGGGTCCTC 1684
                                                                  LENGTH:
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                                                                    4661
                                                                                                                PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 3846;
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Query Match

100.0%;

Score

27;

DB 10;

Length 4661;

Best Local Matches

27; Conservative (

0;

US-09-755-100-2

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APPLICANT: Craig Rosen,
APPLICANT: Steve Ruben
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
FILE REFERENCE: PA101
CURRENT APPLICATION NUMBER: U$/09/925,300
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05988
PRIOR APPLICATION NUMBER: PCT/US00/05988
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 1909-03-12
NUMBER OF SEQ ID NOS: 1890
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 153
LENGTH: 1678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/09755100

Patent No. US20020099189A1

GENERAL INFORMATION:
APPLICANY: SAVITEKY, Kinneret et al.
TITLE OF INVENTION: NOVEL NUCLEIC ACID AND AMINO ACID SEQUENCES
FILE REFERENCE: 2786-0156P

CURRENT FILING DATE: US/09/755,100

CURRENT FILING DATE: 1201-01-08
PRIOR APPLICATION NUMBER: II 128587
PRIOR APPLICATION NUMBER: II 128587
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-08-11
SOUTWARE: PATENT 1999-08-11
SOUTWARE: PATENT 1999-08-11
SEQ ID NO 2
LENGTH: 4661
TYPE: DNA
                                                                                                                                                                            PEATURE:
NAME/REY: misc feature
LOCATION: (163)
OTHER INFORMATION: n equals a,t,g, or c
US-09-925-300-153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 100.0%; Score 27; DB 10; Length 4661; Best Local Similarity 100.0%; Pred. No. 0.0096; Matches 27; Conservative 0; Mismatches 0; Indels 0;
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                                                                     Score 23.8; DB 10;
Pred. No. 0.19;
D; Mismatches 2;
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Patent No. US20020110821A1
REMERAL INFORMATION:
APPLICANT: Ebner, Reinhard
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Usin
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-74
CURRENT APPLICATION NUMBER: US/09/962,832
CURRENT FILING DATE: 2001-09-25
PRIOR APPLICATION NUMBER: US/60/235,077
PRIOR RILING DATE: 2000-99-25
PRIOR RILING DATE: 2000-99-25
NUMBER OF SEQ ID NOS: 259
SOFTWARE: Patentin version 3.0
SEQ ID NO 101
LENGTH: 871
TYPE: DNA
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GENERAL INFORMATION:

APPLICANT: ROSEN et al.

FILE REFERENCE: PAIO5

FULE REFERENCE: PAIO5

CURRENT APPLICATION NUMBER: US/09/925,297

CURRENT FILING DATE: 2001-08-10

PRIOR APPLICATION NUMBER: PCT/US00/05989

PRIOR APPLICATION NUMBER: POI/124,270

PRIOR APPLICATION NUMBER: 00/124,270

PRIOR APPLICATION NUMBER: 
RESULT 15
US-09-880-107-2307
; Sequence 2307, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
; APPLICANT: Horne, Darci T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
US-09-925-297-286
"""Ance 286, Ap
                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-962-832-101
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US-09-962-832-101
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                                                                                                                                                                                                                      194 TCCTGGTGCACCGCCAGTGGGTGCTC 219
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23; Conservative
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US20020081659A1
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88.5%;
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Pred. No. 2.3;
0; Mismatches
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RESULT 12 US-09-925-300-153

Sequence 153, Application US/09925300 Patent No. US20020151681A1 GENERAL INFORMATION:

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; LENGTH: 4661 ; TYPE: DNA ; ORGANISM: Homo sapiens US-09-755-100-2

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1 GTTCTGGTGCACCCCAGTGGGTCCTC 27

Query Match

ocal

Similarity 92.0 25; Conservative

88.1%; 92.6%;

0;

TYPE: DNA
ORGANISM: Homo sapiens

Matches

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APPLICANT: Vockley, Joseph G.
APPLICANT: Scherf, Uwe
APPLICANT: Gene Logic, Inc.
TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
FILE REFERENCE: 44921-5028-W0
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: US 60/211,379
PRIOR PELICATION NUMBER: US 60/211,379
PRIOR PELICATION NUMBER: US 60/237,054
PRIOR APPLICATION NUMBER: US 60/237,054
PRIOR PELICATION NUMBER: US 60/237,054
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Maximum Match 100%
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Maximum DB seq length: 2000000000
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Perfect score:
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         N_Geneseq_101002:*
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2: /SIDS2/gcgdata/g
3: /SIDS2/gcgdata/g
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6: /SIDS2/gcgdata/g
7: /SIDS2/gcgdata/g
8: /SIDS2/gcgdata/g
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Gapop 10.0 , Gapext 1.0
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3: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA199DAT:*

4: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA199DAT:*

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10: /SIDS2/gcgdata/geneseq/geneseqn-emb1/NA109DAT:*

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AAT90073
AAT90074
ABV38513
ABV1435
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ABV07463
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ALIGNMENTS

AAD21363 standard;

DNA; 27 ВP

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AAD21363
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DT 28-J
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ET CDS
FT C Unidentified. 10-APR-2001; WO200176622-A2 Prostate-specific antigen; PSA; immunogenic peptide; immune response; therapy; cancer; prostate cancer; tumour metastasis; prophylaxis; ds. Prostate specific antigen (PSA) derived peptide, CLP316 encoding DNA 28-JAN-2002 AAD21363; (AVET) AVENTIS PASTEUR LTD 10-APR-2000; 2000US-195456P 18-OCT-2001. 2001WO-CA00473 (first entry) /*tag= a
/product= "PSA derived peptide, CLP316"
/note= "CDS does not include start and stop codon" Location/Qualifiers 1..27 /partial

P-PSDB; AAE13025.

2001-663015/76.

Pedyczak A,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The patent discloses immunogenic peptides of prostate-specific antigen (PSA) and nucleic acids encoding them. The peptides of the invention are used to prepare a medicament to elicit an immune response in an animal. They are used to treat cancer such as prostate cancer and tumour metastasis. They are also useful for prophylaxis, for preparing monoclonal or polyclonal antibodies, and in conventional techniques of immunology, molecular biology, cell biology and recombinant DNA technology. The present sequence is a DNA encoding that are the present sequence is a DNA encoding that are the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the present sequence is a DNA encoding that the pr
                    Prostate specific antigen multiple epitope peptide - useful i vaccine to produce immune response against prostate specific in prostatic tumour cell
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                                                                                                                                                                              Schlom J,
                                                                                                                                                                                                                                                                                           20-MAR-1996;
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DB; AAW27387.
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27; Conser
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DEPT HEALTH & HUMAN SERVICES
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llarity 100.0%;
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                                                                                                                                                                              Zaremba
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Pred. No. 0.0051;
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RESULT 3
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                                                                                                                             Prostate specific antigen multiple epitope peptide - useful in vaccine to produce immune response against prostate specific antigen in prostatic tumour cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prostate specific antigen; PSA; epitope; vaccine; immune respon
prostatic tumour cell; human leukocyte antigen; HLA; cytotoxic;
T lymphocyte; cancer; ds.
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                                                                                                  Claim 23; Page 67; 85pp; English.
                                                                                                                                                                                          WPI; 1997-480226/44.
                                                                                                                                                                                                                      Schlom J,
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(USSH ) US DEPT HEALTH & HUMAN SERVICES.
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27; Conser
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Pred. No.
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The present sequence encodes a prostate specific antigen oligo-epitope peptide (PSA-OP) for use in a virus vector of the present invention. The PSA-OP comprises more than 1 adjoined PSA-OP which generates a prostate specific response in a portion of the human population. The PSA-OP can be used to produce PSA specific cytotoxic T lymphocytes; in vivo or in vitro, which can kill or prevent the establishment or growth of PSA

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                                     a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a therapy for inhibiting prostate can a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          positive tumour cells, i.e. to prevent or treat prostatic cancer. The products may be used to elicit a PSA specific immune response. When a recombinant virus is used to elicit the response it is coadministered with a carrier, a biological response modifier e.g. interleukin 2, 6 o 12, interferon, tumour necrosis factor, granulocyte/macrophage-colony stimulating factor or cyclophosphamide and an adjuvant selected from RIBI Detox, OS21, alum and incomplete Freund's adjuvant pSA-op comprises multiple PSA epitopes, allowing an immune response to PSA in a population of humans having more than 1 HLA class I molecule type to
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                                                                                                                                                                                                                                                                    Claim 1;
                                                                                                                                                                                                                                                                                                           Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, uses for detecting presence of prostate cancer, stage of prostate cancer
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the prostate of
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Pred. No. (
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16-MAR-2000;
25-MAY-2000;
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                                                          (d) assessing the efficacy of the concert in a patient; in a patient; in a patient; (e) selecting a composition for inhibiting prostate cancer in (f) assessing the prostate cell carcinogenic potential of a (g) determining whether prostate cancer has metastasized in (g) determining the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence of prostate cancer the aggressiveness or indolence or aggressiveness or indolence the aggressiveness or indolence the aggressiveness or indolence the a
                                                                                                                                                                                                                            The invention relates to an isolated nucleic acid molecule (I) com a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for: (a) assessing whether a patient is afflicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                    (C) assessing the efficacy of a therapy for
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(h) assessing the aggressiveness or indolence of prostate cancer in
patient;(I) is also useful as a pharmacodyanamic or pharmacogenomic marker.
                                      patient;
(I) is also useful
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2000US-189862P.
2000US-207454P.
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16-MAR-2000; 2000US-189862P.
25-MAY-2000; 2000US-207464P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-25281P.
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(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
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(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) selecting a composition for inhibiting prostate cancer in a patient (e) assessing the prostate cell carcinogenic potential of a compound;
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                                                           Sequence 466 BP; 98 A; 140 C; 136 G; 92 T; 0 other;
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27; Conserv
                                                                                      also useful as a pharmacodyanamic or pharmacogenomic marker
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              Score 27;
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RESULT 7
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16-MAR-2000; 2000US-1898629-
25-MAY-2000; 2000US-207454P.
09-JUN-2000; 2000US-211314P.
18-JUL-2000; 2000US-219007P.
13-DEC-2000; 2000US-255281P.
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(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) selecting a composition for inhibiting prostate cancer in a patient;
(f) assessing the prostate cell carcinogenic potential of a compound;
(g) determining whether prostate cancer has metastasized in a patient;
(h) assessing the aggressiveness or indolence of prostate cancer in a patient;
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pharmacogenomic marker;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 328 AAACTTCAGTGTGTGGACCTCCATGTT
   286
is also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                 Ch 100.0%; Score 27; Di Similarity 100.0%; Pred. No. 0.027; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endege WO,
                                                                                                                                                                                                                               103 A; 138 C; 136 G; 99 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cytostatic; carcinogen; pharmacodyanamic gene; ss.
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                                                        27
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دو
مدد م<del>و</del> د حدد
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                                                                                                                                               0.0086
                                                                                                                                                                             23;
                                                                                                                                                                          Length 476;
                                                                                                                        Indels
                                                                                                                    0;
                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tate of
useful
cer -
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RESULT 9
ABV44248
ID ABV4
XX
AC ABV4
XX
DT 16-S
XX
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                                                                                                                                                                        Query
Best I
                                                                                                                                                              Matches
                                                                                                                                                                                                                                        cancer in a patient;
(d) assessing the eff
in a patient;
(e) selecting a compo
(f) assessing the pro
(g) determining wheth
(h) assessing the agg
patient;
(T) is a like work.
                                                                                                                                                                                                                                                                                                                                   The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 1204-1205; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer, stage of prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-FEB-2000;
16-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV07463;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-MAY-2000;
09-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200160860-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABV07463 standard; cDNA; 492
           16-SEP-2002
                                 ABV44248;
                                                       ABV44248
                                                                                                                                                                                                          Sequence 492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schlegel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MILL-) MILLENNIUM PREDICTIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-JUL-2000;
13-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-FEB-2001;
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                                                                                                                245
                                                                                                                                                                          Local
                                                                                                                                       Н
                                                                                                                                                                                                                                                    selecting a composition for inhibiting prostate cancer in a patient; assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                       AAACTTCAGTGTGTGGACCTCCATGTT 27
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                                                                                                                AAACTTCAGTGTGTGGACCTCCATGTT 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                prostate
                                                                                                                                                                                                                                  also
                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ŗ,
                                                        standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; 2000US-183319P.
; 2000US-189862P.
; 2000US-207454P.
; 2000US-211314P.
; 2000US-219007P.
; 2000US-255281P.
                                                                                                                                                                                                                                  useful as
                                                                                                                                                              Conservative
                                                                                                                                                                                                           BP; 102 A; 141 C; 115 G;
          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Endege
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression marker cDNA 7454.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               marker;
                                                                                                                                                                                                                                                                                                               efficacy of a therapy
                                                        CDNA;
          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
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                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                 a pharmacodyanamic or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic;
gene; ss.
                                                        616
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                                                        ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDICINE
                                                                                                                                                                        Pred.
                                                                                                                                                                                    Score 27;
                                                                                                                                                              Mismatches
                                                                                                                                                                           v
Ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          carcinogen;
                                                                                                                          .:
                                                                                                                                                                                                           134 T; 0 other;
                                                                                                                                                                                                                                                                                                               for inhibiting prostate cancer
                                                                                                                                                                                    DB 23;
                                                                                                                                                                          . 0086;
                                                                                                                                                                                                                                pharmacogenomic
                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pharmacodyanamic
                                                                                                                                                                                  Length 492;
                                                                                                                                                              0
                                                                                                                                                              Gaps
                                                                                                                                                              0
                                                                                                                                      RESULT 10
ABV37397/c
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                                                                                                                                                                                                        QΥ
                                                                                                                                                                                                                                Query Match
Best Local S
Matches 27
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(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                                                                                                                                                                                                                                               cancer in a patient; (d) assessing the ef
                    Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                            Sequence
Homo sapiens
                                                                             16-SEP-2002
                                                                                                     ABV37397;
                                                                                                                           ABV37397 standard; cDNA; 632
                                                                                                                                                                                                                                                                                                                                    (e) selecting a composition for inhibiting prostate cancer in a patient;(f) assessing the prostate cell carcinogenic potential of a compound;(g) determining whether prostate cancer has metastasized in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schlegel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-MAR-2000;
25-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pharmacogenomic
                                                                                                                                                                                                                                                                                                                         (g) determinin
(h) assessing
                                                                                                                                                                                                                                                                                                                                                                       in a patient;
                                                                                                                                                                                    252
                                                                                                                                                                                                           Н
                                                                                                                                                                                                                                                                                                    is also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                   AAACTTCAGTGTGTGGACCTCCATGTT
                                                                                                                                                                                                           AAACTTCAGTGTGTGGACCTCCATGTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001-662795/76.
                                                       prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prostate
                                                                                                                                                                                                                                 . Similarity
27; Conserv
                                                                                                                                                                                                                                                                              616
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2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-2119007P.
2000US-255281P.
                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                             BP;
                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Endege
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001WO-US05171
                                                                                                                                                                                                                                                                                                                         the aggressiveness
                                                       expression
                    marker; gene; ss
                                cancer; cytostatic;
                                                                                                                                                                                                                                                                              122
                                                                                                                                                                                                                                                                                                                                                                                  efficacy of a therapy for inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO,
                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                            A; 176
                                                        marker
                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                              Ç
                                                                                                                                                                                                                                Score 27; DB 23
Pred. No. 0.009;
D; Mismatches
                                                                                                                                                                                                                                                                            186 G;
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278 27

CDNA

37388

carcinogen;

pharmacodyanamic

marker;

or

indolence of

prostate

cancer in

a

prostate cancer

) comprising of the

132 T; 0 other;

23; Ò,

Length

616;

0;

Gaps

0

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Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, used for detecting presence of prostate cancer, stage of prostate cancer
Claim 1; Page 8788; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human prostate expression marker cDNA 44239
                                                                                                                                                                                                                              (MILL-) MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer; cytostatic; carcinogen;
marker; gene; ss.
                                                                                                                                                                                      Monahan
                                                                                                                                                                                      JE;
                                                                                                                                                                                                                                 INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;-
_-:
                                                                                                                                                                                                                                                                                                                                   pharmacodyanamic marker;
                                                    tate of
useful
cer -
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RESULT 11
AAT92976
ID AAT92
XX AAT92
AC AAT92
XX 24-AH
XX CDNA
XX D1agr
KW D1agr
KW D506e
XX H0mo
XX H0mo
XX W0973
XX 23-OC
XX 23-OC
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                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 27
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16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for:

(a) assessing whether a patient is afflicted with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting prostate cancer in a patient;
(e) assessing the prostate cancer in a patient;
(f) assessing the prostate cell carcinogenic potential of a compound;
(g) determining whether prostate cancer has metastasized in a patient;
(h) assessing the aggressiveness or indolence of prostate cancer in a patient.
                                                                              Diagnosis; prostate cancer; prostate specific antigen; PSA; metastatic-prostate cancer; prostate cell; secondary tumour; bone metastatic anchoring; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer -
                                                                                                                             cDNA for prostate specific antigen (PSA).
                                                                                                                                                                                                                                                                                                                                                      Sequence 632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 7678; 11750pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-662795/76.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2001; 2001WO-US05171
            23-OCT-1997
                                 WO9739139-A1
                                                                                                                                                      24-APR-1998
                                                                                                                                                                                                   AAT92976 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200160860-A2.
                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                         patient;(I) is also useful as a pharmacodyanamic or pharmacogenomic marker.
                                                                                                                                                                                                                                                           326
                                                                                                                                                                                                                                                         ch 100.0%; Score 27; DB 1 Similarity 100.0%; Pred. No. 0.00 27; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ₽,
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2000US-211314P.
2000US-219007P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Endege WO,
                                                                                                                                                                                                                                                                                                                                                   BP; 130 A; 182 C; 160 G; 160 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2000US-183319P
2000US-189862P
                                                                                                                                                   (first entry)
                                                                                                                                                                                                    639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Monahan JE;
                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                           300
                                                                                                                                                                                                                                                                                                                DB 23; Length 632; .009;
                                                                                                                                                                                                                                                                                                        0;
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                                                                                                                                                                                                                                                                                                        0;
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ABV22297
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                                                                                                                                                                                                                                    Homo
                                   Schlegel
                                                        (MILL-)
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Query Match
                          17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
13-DEC-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CDNA sequences AAT92974-77 encode the prostate specific antigen (PSA). The detection of PSA mRNA in peripheral'blood is associated with metastatic prostate cancer. Detection of PSA mRNA in the bloodstream indicates that prostate cells are circulating in the blood and confirms diagnosis of cancer. Circulating prostate cells also indicate a risk of a secondary tumour or bone metastatic anchoring. Detection of the PSA mRNA after prostatectomy or radiotherapy indicates a risk that the prostate cancer has spread and that the surgery and radiotherapy was not effective. PCR primers AAT92971-73 were used for the monitoring and diagnosis of patients with prostate cancer. The method comprises the detection of PSA mRNA by reverse transcriptase polymerase chain reaction
                                                                                                                                                                                                                                                                                                   Human prostate expression marker cDNA 22288
                                                                                                                                                                                                                                                                                                                                   13-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                           ABV22297
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16-APR-1997;
                                                                                                                                                                  23-AUG-2001.
                                                                                                                                                                                              WO200160860-A2
                                                                                                                                                                                                                                                        Human; prostate pharmacogenomic
                                                                                                                                                                                                                                                                                                                                                                ABV22297;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Monitoring or diagnosis of prostate oppositate-specific antigen mRNA using
                                                                                                                                    20-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RT-PCR).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    486 AAACTTCAGTGTGTGGACCTCCATGTT 512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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                                                                                                                                                                                                                          sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               l Similarity
27; Conserv
MILLENNIUM PREDICTIVE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                           standard; cDNA; 651
                          2000US-183319P.
2000US-189862P.
2000US-207454P.
2000US-211314P.
2000US-2119007P.
2000US-255281P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP;
                                                                                                                                    2001WO-US05171
                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                      marker; gene; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             133 A; 191 C; 185 G; 130 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 27;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                      carcinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer - |
g specific
                                                                                                                                                                                                                     · [1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 18;
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primers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                       pharmacodyanamic marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 639;
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RESULT 13
ABV28130
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of t specification or its complement. (I) is useful for:
(a) assessing whether a patient is affiliated with prostate cancer;
(b) monitoring the progression of prostate cancer in a patient;
(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                            17-FEB-2000;
16-MAR-2000;
25-MAY-2000;
09-JUN-2000;
18-JUL-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Page 3854; 11750pp; English.
                                                                                                                                                                                                                                                                                                                           Human; prostate pharmacogenomic
                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (c) assessing the efficacy of a test compound to inhibition cancer in a patient;
(d) assessing the efficacy of a therapy for inhibiting in a patient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated nucleic acid molecule associated with cancerous state of prostate cells and correlating with presence of prostate cancer, useful for detecting presence of prostate cancer -
          Claim
                                  ior
                                           prostate
                                                      Novel isolated
                                                                                                     Schlegel
                                                                                                                           (MILL-)
                                                                                                                                                                                                                                 20-FEB-2001;
                                                                                                                                                                                                                                                         23-AUG-2001.
                                                                                                                                                                                                                                                                             WO200160860-A2
                                                                                                                                                                                                                                                                                                      Homo
                                                                                                                                                                                                                                                                                                                                                              Human prostate
                                                                                                                                                                                                                                                                                                                                                                                      16-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                              ABV28130
                                                                                                                                                                                                                                                                                                                                                                                                                                  ABV28130 standard; cDNA; 651
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 selecting a composition for inhibiting prostate cancer in a patien assessing the prostate cell carcinogenic potential of a compound; determining whether prostate cancer has metastasized in a patient; assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention relates
                               detecting
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                                                                              2001-662795/76
          1;
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                                                                                                                           MILLENNIUM PREDICTIVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      651
                                                                                                    70
         Page
                                             cells
                                                                                                                                                ; 2000US-183319P.
; 2000US-189862P.
; 2000US-207454P.
; 2000US-211314P.
; 2000US-2119007P.
; 2000US-255281P.
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                                                                                                                                                                                                                                 2001WO-US05171.
                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     B₽;
                               presence of prostate
          5827;
                                         nucleic acid molecule associated with cancerous state and correlating with presence of prostate cancer, use
                                                                                                                                                                                                                                                                                                                                                              expression marker cDNA 28121.
                                                                                                                                                                                                                                                                                                                             cancer;
marker;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    164 A; 171 C;
         11750pp; English
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                                                                                                                                                                                                                                                                                                                                       cytostatic;
                                                                                                     Monahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 isolated nucleic acid molecule (I) comprising in Tables 1-9 (ABV00010-ABV62213) of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 27;
Pred. No.
                                                                                                                           MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     156
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                                cancer,
                                                                                                                                                                                                                                                                                                                                       carcinogen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     160 T;
                                                                                                                            INC
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                                stage
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                                                                                                                                                                                                                                                                                                                                       pharmacodyanamic
                                of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 651;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                prostate cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            marker
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                                                                                                                                                                                                                                                                                                                                        marker;
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RESULT 14
AAX08947
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Best Local
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                                                                                                                                                                                       (KLEE/)
(MAYO-)
(TIND/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to an isolated nucleic acid molecule (I) come a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is specification or its complement. (I) is sefful for:

(a) assessing whether a patient is afflicted with prostate cancer;

(b) monitoring the progression of prostate cancer in a patient;

(c) assessing the efficacy of a test compound to inhibit prostate
                                                                                                                                                     Klee
                                                                                                                                                                                                                                                                                                                                                                                                         Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX08947;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer in a patient;
(d) assessing the efficacy of a therapy for in a patient;
Human kallikrein 2 (hK2) is expressed at elevated levels relative the prostate cancer antigen, prostate specific antigen (PSA) by breast cancer cells when stimulated by androgens. Detecting level
                                                                            the
                                                                                                               P-PSDB;
                                                                                                                                                                                                                                                                                                                  WO9859073-A1.
                                                                                                                                                                                                                                                                                                                                                                   CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human prostate specific antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX08947
                                                   Disclosure;
                                                                                                                                                                                                                                          20-JUN-1997;
                                                                                                                                                                                                                                                                  19-JUN-1998;
                                                                                                                                                                                                                                                                                           30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                               Human kallikrein 2; hK2; breast cancer; diagnosis; dete treatment; monitoring; prostate specific antigen; PSA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-APR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (e) selecting a composition for inhibiting prostate cancer in a patient;(f) assessing the prostate cell carcinogenic potential of a compound;(g) determining whether prostate cancer has metastasized in a patient;(h) assessing the aggressiveness or indolence of prostate cancer in a
                                                                                                                                                                             (\nnox)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (I) is also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           249
                                                                         of human kallikrein 2 - as a marker 1 diagnosis, prognosis, monitoring and
                                                                                                                1999-105632/09
DB; AAW96187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAACTTCAGTGTGTGGACCTCCATGTT 275
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                                                                                                                                                     9
                                                                                                                                                                                                                                                                                                                                                                                                        sapiens
                                                                                                                                                                       ) KLEE G G.
) MAYO FOUNDATION.
) TINDALL D J.
') YOUNG C Y F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                     Tindall
                                                 Page 34-35; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                           /label=
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                                                                                                                                                                                                                                                                                                                                                                      Cocation/Qualifiers
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                                                                                                                                                  Young
                                                                                                                                                                                                                                                                                                                                  a
Prostate_specific_antigen
                                                                          as a marker for developing products for 
onitoring and treatment of breast cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                     CYF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 27;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .0091;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tate cancer;
patient;
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; prostate specific antigen; PSA; cytostatic; antiviral; immunostimulant; cell-mediated immune response; tumour; breast cancer; virus infection; prostate cancer; colorectal cancer; pancreatic cancer; lymphoma; leukaemia; hepadnavirus; lentivirus; herpesvirus; human immunodeficiency virus; HIV; flavivirus; pestivirus; gene; ss.
   Inducing a cell-mediated immune response against a target antigen, reducing undesired cells and stimulating presentation of an antigen a cell, comprises administering a polynucleotide encoding a variant an antigen -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 711 BP; 148 A; 208 C; 206 G; 149 T; 0 other;
an antigen
                                                                                                                                         WPI; 2002-527524/56.
P-PSDB; AAU98921.
                                                                                                                                                                                                                                                                               (AMBI-) AMERICAN FOUND BIOLOGICAL RES (MINC) MINCHEFF M S. (LOUK) LOUKINGV D I. (ZOUB/) ZOUBAK S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-2001; 2001WO-US45626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200240059-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cDNA encoding human prostate specific antigen (PSA) variant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK86205
                                                                                                                                                                                                                            Mincheff MS,
                                                                                                                                                                                                                                                                                                                                                                                                                            01-NOV-2000; 2000US-0704232.
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27; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 llarity 100.0%; Score 27; DB 20; Length 711; 100.0%; Pred. No. 0.0092; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                            Loukinov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
7..744
/*tag- a
/product= "Prostate specific antigen (PSA) variant
with signal sequence deleted"
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The invention

6

a method of inducing

Ø

cell-mediated

Disclosure; Page 121-122; 146pp; English.

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response against a cell comprising a target antigen (I) in a subject, treating a subject having undesired cells, for example tumour cells or virally infected cells (C), reducing the number of (C) in a subject and stimulating presentation of (I) by a cell. This is done by administering a polynucleotide (II) encoding a variant of (I), so that (II) expressed in a cell and cell-mediated immune response is induced. The method can be used to treat prostate cancer, breast cancer, colorectal cancer and pancreatic cancer, as well as lymphomas and leukaemias. The method is also useful in treating chronic viral infections such as those caused by hepadnaviruses, lentiviruses (including human immunodeficiency virus (HIV)), herpesviruses and the flaviviruses and pestiviruses. The present sequence represents the cod sequence of human prostate specific antigen (PSA) variant which has the signal sequence deleted, used as a target antigen in the methon of the invention
                                                     of the invention.
                                                                                       which
in the method
                                                                                                                                                                                                                                                                                                                                                                                                     a subject,
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Sequence 744 BP; 153 A; 222 C; 213 G; 156 T; 0 other;

οy Query Match
Best Local Similarity
Matches 27; Conserv 1 AMACTICAGIGIGIGACCICCAIGIT 27 AAACTTCAGTGTGTGGACCTCCATGTT 492 100.0%; Score 27; DE ilarity 100.0%; Pred. No. 0.0 Conservative 0; Mismatches DB 24; Length 744; 0.0093; hes 0; Indels 0; Gaps

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Search completed: November 19, Job time: 191.769 secs 2002, 03:06:38

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Result
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic search, using sw model
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27
1 aaacttcagtgtgta
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1: /cgn2_6/ptodata/2,
2: /cgn2_6/ptodata/2,
3: /cgn2_6/ptodata/2,
4: /cgn2_6/ptodata/2,
5: /cgn2_6/ptodata/2,
6: /cgn2_6/ptodata/2,
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length: 2000000000
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Listing first 45 summaries
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/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/2/ina/backfiles1.seq:*
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  DВ
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                    US-09-100-264-4
US-08-358-7820-13
US-08-605-785-616
US-08-605-785-616
US-08-64-527A-14
US-08-64-527A-14
US-08-64-527A-14
US-08-62-615-7820-1
US-08-596-993-1
US-08-596-993-1
US-08-596-993-7
US-08-767-820A-7
US-08-767-820A-7
US-08-768-622-046B-2
US-08-768-768-99-7
US-08-768-628-768-99-7
US-08-768-89A-7
US-08-768-89A-9
US-08-768-89A-9
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US-08-622-046B-6
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                                                                         GENERAL INFORMATION:
APPLICANT: Croce, Carlo
APPLICANT: Croce, Leonard
APPLICANT: Mulholland, S. Grant
APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Det
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                 Sequence 13, Application US/08358782D Patent No. 5674682
  COMPUTER READABLE FORM
                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock
          STREET: One Liberty CITY: Philadelphia STATE: PA ZIP: 19103
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E: Woodcock Washburn Kurtz Mackiewicz One Liberty Place 46th. Floor

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Detecting

Micrometastasis

of Prostate

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larity Conservat			444444444477777
100 100 rative	4, Application US/09100 6.6235486 NEORMATION: TI:Young, Charles Y.F. TI: Tindall, Donald J. TI: Klee, George G. INVENTION Method for INVENTION NUMBER: US/ PILING DATE: 1998-06-19 APPLICATION NUMBER: US/ FILING DATE: 1997-06-20 F SEQ ID NOS: 15 F SET IN NOS: 1		832 832 832 832 832 832 1341 13541 13541 13541 13541 2417 2417 2417 2417 2360 3360 3360
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MEDIUM TYPE: COMPUTER: IF

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; MOLECULE TYPE: DNA (genomic); HYPOTHETICAL: NO; ANTI-SENSE: NO US-08-358-782D-13
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TELEPHONE: 215-568-3100
TELEPAX: 215-568-3439
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 992 base pairs
TYPE: nucleic acid
STRANDEDNESS: Single
TOPOLOGY: 11 near
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Best Local S
Matches 27
PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/358,782

FILING DATE: 15-DEC-1994

ATTORNEY/AGENT INFORMATION:

NAME: Beardell, Lori Y.

REGISTRATION NUMBER: 34,293

REFERENCE/DOCKET NUMBER: TJU-1327

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATE PATENTIA PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
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APPLICANT:
APPLICANT:
APPLICANT:
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NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-1327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PATENTIN RECURRENT APPLICATION NUMBER: UNFILING DATE: 15-DEC-1
                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis
TITLE OF INVENTION: Prostate
NUMBER OF SEQUENCES: 14
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IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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N: 435
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APPLICANT: MEDIET, WILLIAM

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER

TITLE REFERENCE: 210121.427C16

CURRENT APPLICATION NUMBER: US/09/605,785

CURRENT APPLICATION NUMBER: US/09/605,785

CURRENT FILING DATE: 2000-06-27

NUMBER OF SEQ ID NOS: 835

SOFTWARE: FASTSEQ FOR WINDOWS VETSION*3.0

SEQ ID NO 616

SEQ ID NO 616

LENGTH: 1350

TYPE: DNA

ORGANISM: Homo sapien

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; HYPOTHETICAL: I
ANTI-SENSE: NO
US-08-764-527A-13
                                                               US-08-358-782D-14
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              GENERAL (
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Patent N
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                                                                                                                                                                                 Local Similarity es 27; Conserv
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              NFORMATION:
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                                               Application US/08358782D
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Gomella, Leonard Mulholland, S. Grant

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                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
NAME: Beardell, Lori Y.
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MEDIUM TYPE: Floppy disk
                                    CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                        STATE: 19103
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ITTLE OF INVENTION: Methods of Detecting Micrometastasis ITTLE OF INVENTION: Prostate
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FILING DATE: 15-DEC
CLASSIFICATION: 435
                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
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   APPLICATION NUMBER:
                                                                        SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
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DEDNESS: single
                                                                                                                                                                                                                                                                                                               Philadelphia
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                                                                                                                                                                                                                                                                                                                                            E: Woodcock Washburn Kurtz Mackiewicz & One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1462 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mulholland, S. Grant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Moreno, Jose
                                                                 PatentIn Release #1.0, Version #1.25
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SYSTEM: PC-DOS/MS-DOS
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US/08/764,527A
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Best Local Similarity
Watches 27; Conserve
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US-08-764-527A-14
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Patent No. 5840494
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                                         SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                    APPLICATION NUMBER: 08/2 FILING DATE: 15-APR-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 1462 base pairs
TYPE: nucleic acid
                                                                                                            TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                PRIOR APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS:
                                                                                                                                          NAME: White, John P
REGISTRATION NUMBER:
                                                                                                                                                                                                                                          APPLICATION NUMBER: FILING DATE: 18-APP
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                                                                                 TELEFAX:
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                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                    : New York
E: New York
TRY: U.S.A.
           nucleic acid
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                                                                    NE: (212) 977-95
422523 COOP UI
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                                                                                                          NUMBER: 28,678
NUMBER: 43677/JPW/TEP
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18-APR-1997
V: /эг
single
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                                                                                                                                                                                                 08/229,391
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Matches 27; Conser
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                                                                                                                                   Query Match
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, 1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/718,547
FILING DATE:
                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: SEQUENCE: 1729 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 4367
TELECOMMUNICATION INFORMATION:
TELEPHANE: (212) 278-0400
TELEFAX: (212) 391-0525
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                                                                                                                                                                                                                                                          MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS
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TITLE OF INVENTION: A MC
TITLE OF INVENTION: Prov
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                                  813 AAACTTCAGTGTGTGGACCTCCATGTT 839
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les 27; Conser
                                                                                                                                                                                                      NAME/KEY:
                                                                                                                                                                                                                                        ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
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STATE: New York
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                                                                1 AAACTTCAGTGTGTGGACCTCCATGTT 27
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1185 Avenue of the Americas
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                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Katz, Aaron E.; Buttyan, Ralph; Raffo, Anthony;
                                                                                                                                                                                                                                       Homo sapiens
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                                                                                                                                                                                                                                                                                                         single
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                                                                                                                100.0%;
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                                                                                                                  Score 27;
Pred. No.
                                                                                                     Mismatches
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US-09-056-105-31
; Sequence 31, A
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US-08-294-611-1/c
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                                                                                    TELEFAX: (215) 568-3439 INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
FILING DATE:
                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                               REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TUTELECOMMUNICATION INFORMATION
                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Methods of Detec
TITLE OF INVENTION: Micrometastasis
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: WORDPERFECT 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: One Libert;
CITY: Philadelphia
STATE: PA
                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: UZIP: 19103
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                                                                                                                TELEPHONE:
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es 27; Conserv
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Nucleic
)EDNESS: Single
)GY: Linear
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                                                                                                                  (215)
                                                                                                                                                                                 Beardel
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Micrometastasis Of Prostate Cancer
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TJU-0722

GENERAL INFORMATION:
APPLICANT: KIPPS, THOMAS J.
APPLICANT: WU, YUNQI
TITLE OF INVENTION: VACCINES WITH ENHANCED INTRACELLULAR
TITLE OF INVENTION: PROCESSING
FILE REFERENCE: 233/221 Patent No. 6287569 CURRENT APPLICATION NUMBER: US/09/056,105 CURRENT FILING DATE: 1998-04-06 EARLIER APPLICATION NUMBER: 60/043,467 EARLIER FILING DATE: 1997-04-10 NUMBER OF SEQ ID NOS: 35 4200 AAACTTCAGTGTGGACCTCCATGTT 4226 1 AAACTTCAGTGTGTGGACCTCCATGTT 27 FastSEQ for Windows Version 3.0 100.0%; ilarity 100.0%; Conservative 0 Score 27; DB 4; Pred. No. 0.0018 Mismatches Length 7130; Indels 0;

Mulholland, S.

Grant

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RESULT 12
US-08-596-993-1/c
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US-08-358-782D-1/c
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US-08-294-611-1
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                                       Sequence 1, Applia Patent No. 568864
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                                                                                                                                                                              Matches
                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
GENERAL INFORMATION:
APPLICANT: Croce, Carlo
APPLICANT: Gomella, Leonard
                                                                                                                                                                                                                                                                                                                                            TELEFAX: 215-568-3439
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/358
FILING DATE: 15-DEC-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Croce, Carlo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                  MOLECULE TYPE: DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of Detecting Micrometastasis of Prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: One Liberty
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                          20
                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                        2 AACTTCAGTGTGTGGACCTC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                 ENGTH:
                                                                                                                         AACTTCAGTGTGGACCTC 1
                                                                                                                                                                               20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19103
                                                                                                                                                                                                                                                                                                                  nucleic acid
                                                       Application US/08596993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        E: Woodcock Washburn Kurtz Mackiewicz & One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                        215-568-3100
215-568-3439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mulholland, S. Grant
                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Moreno, Jose
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ő
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Floppy disk
                                                                                                                                                                                                                                                                            DNA (genomic)
                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            74.18;
                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leonard
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                                                                                                                                                                               0;
                                                                                                                                                                                             Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                   TJU-1327
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Pred. No.
                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                             1.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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2:
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                                                                                                                                                                                                       Length 20
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                                                                                                                                                                               Indels
                                                                                                                                                                              0;
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                                                                                                                                                                               Gaps
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Query Match
Best Local Similarity
Watches 20; Conserv?
                                                                                                                                                                                                                                                                               RESULT 13
US-08-764-527A-1/c
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US-08-596-993-1
                                                                                                                                                                                                                                               Sequence 1, Application US/08764527A Patent No. 5939258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/294,611
FILING DATE: 23-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/973,322
FILING DATE: 29-OCT-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (215) 568-34
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Mulhollar APPLICANT: Moreno, J APPLICANT: Fischer, TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: TJ
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                 NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                   APPLICANT: Moreno, Jose
APPLICANT: Fischer, Rainer
TITLE OF INVENTION: Methods of
TITLE OF INVENTION: Prostate
                                                                                                                                                                            APPLICANT: Croce, Carlo
APPLICANT: Gomella, Leonard
APPLICANT: Mulholland, S. G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
           STREET: One L.CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
STATE: 19103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM COMPONER: OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/596,993 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Mackiewicz & No.
                                                                                                                                                                                                                                                                                                                                                                          AACTTCAGTGTGTGGACCTC 21
                                                                                                                                                                                                                                INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mark DeLuca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic
                                                      E: Woodcock Washburn Kurtz Mackiewicz
One Liberty Place 46th. Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: Mackiewicz & No. 5688649ris
One Liberty Place - 46th Floor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fischer, Rainer
VENTION: Methods of Detecting
VENTION: Micrometastasis Of Prostate Cancer
                                                                                                                                                                                                                                                                                                                                                                                                         74.1%;
llarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Moreno, Jose G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM COMPATIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Woodcock Washburn Kurtz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DISKETTE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    568-3439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                  Grant
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 20; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TJU-1657
                                                                                                                                    Detecting
                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                     Micrometastasis
                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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0,

Gaps

0;

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STREET: P.O. Box 2938

CITY: Minneapolis

STATE: M.

COUNTRY: USA

ZIE: 55402

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFFWARE: MS-LCATION DATA:
APPLICATION NUMBER: US/08/622,046B

CLASSIFICATION: 514

ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 30,440

REGISTRATION NUMBER: 30,440

REGISTRATION NUMBER: 30,440

RESERBACE/DOCKET NUMBER: 30,440

TELEPHANE: 612-373-6903

TELEPHANE: 612-373-6903

TELEPHANE: 612-373-9330

TELEPHANE: 612-373-9330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: DNA HYPOTHETICAL: NO ANTI-SENSE: NO US-08-764-527A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US/08/358,782
FILLING DATE: 15-DEC-1994
ATTORNEY/AGENT INFORMATION:
NAME: Beardell, Lori Y.
REGISTRATION NUMBER: 34,293
REFERENCE/DOCKET NUMBER: TJU-1327
TELECHONE: 215-568-3100
TELECHONE: 215-568-3100
INFORMATION FOR SEQ ID NO: 1:
TELEPAX: 215-568-310
INFORMATION FOR SEQ ID NO: 1:
INFORMATION FOR SEQ ID NO: 1:
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/08622046B Patent No. 6103237 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match 74.1%; Score 20; DB 2; Length 20; Best Local Similarity 100.0%; Pred. No. 1.2; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Saedi, Moham
APPLICANT: Mikolajczyk,
TITLE OF INVENTION: Sta
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/764,527A
                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth, P.A. STREET: P.O. Box 2938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mohammed S
                                                                                                                                                                                                                                                                                                                                                                                                                       zyk, Stephen D.
Stable Variant HK2 Polypeptide
21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                               NAME: WOESSNEY WATER D.
REGISTRATION NUMBER: 30,440
REFERENCE/DOKET NUMBER: 476.001US1
RELECOMMUNICATION INFORMATION:
TELEPHAX: 612-33-903
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 711 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: CDNA
US-08-622-046B-13
                                                                                                             В
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Best Local S
Matches 20
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US-08-622-046B-13
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TOPOLOGY: linear
MOLECULE TYPE: CDN
US-08-622-046B-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13. Application US/08622046B
PATENT NO. 6103237
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Mikolajczyk, Stephen D.
TITLE OF INVENTION: Stable Variant
NUMBER OF SEQUENCES: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 67.4%;
Best Local Similarity 87.0%;
Matches 20; Conservative
                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERAPTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/622,046B
FILING DATE: 26-MARCH-1996
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                             completed: November 19, 2002, 05:20:22 ne : 53.9615 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSE: Schwegman, Lundberg, Woessner & Kluth, P.A.
STREET: P.O. Box 2938
CITY: Minnapolis
STATE: MN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STALL
COUNTRY: 55402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       439 CTTCAGTGTGTGAGCCTCCATCT 461
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                                                                                                                                                                       Similarity 87.0
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                                                                                                                                                                                         67.48;
87.08;
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Jozyk, Stephen D.
Stable Variant HK2 Polypeptide
S: 21
                                                                                                                                                                          0;
                                                                                                                                                                                         Score 18.2;
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 18.2; DE Pred. No. 14; 0; Mismatches
                                                                                                                                                                     Mismatches
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0;

Gaps

0;

RESULT 14 US-08-622-046B-2